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CC encoding a Propionibacterium acnes protein. The invention also relates to collowing a propionibacterium acnes protein. The invention also relates to collowing a propionibacterium acnes protein. The invention and to collow immunogenic fragments of P. acnes polypeptides. The invention compasses expression vectors and host cells comprising a polypeptide of the invention; antibodies against polypeptides of the invention; antibodies against polypeptides of the collowing in immune response specific for a P. acnes comprising a polypeptide of the invention; a comprision grotein gr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New Propionibacterium acnes polypeptides and polynucleotides encoding polypeptide, useful for diagnosing, preventing or treating acne vulgas or for stimulating an immune response specific for a P. acnes protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; SEQ ID NO 20252; 1481pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-381789/36.
N-PSDB; ACF64526.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Propionibacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Acne vulgaris; antiseborrhoeic; dermatological; antibacterial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Propionibacterium acnes predicted ORF-encoded polypeptide #20252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-OCT-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CORI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        immunostimulant; immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention relates to an isolated polynucleotide (ACF64435-ACF64733)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wang S, Jen S, 1
Vallieve-Douglass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Skeiky YAW, Persing DH,
ing S, Jen S, Lodes MJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B
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Benson DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jones R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Maisonneuve JL;
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RESULT 39
ABU27913
                     CC the 6213 antition required given in the specification where expression of the nucleic acid inhibits proliferation of a cell. Also included are: (1) a vector comprising a promoter operably linked to the nucleic acid cencoding a polypeptide whose expression is inhibited by the antisense concleic acid; (4) a host cell containing the vector; (3) an isolated containing the polypeptide by the containing the vector; (3) an isolated containing the polypeptide; (5) producing the polypeptide; (6) inhibiting cellular containing the polypeptide; (6) inhibiting cellular proliferation; (7) identifying a compound that influences the activity of the gene product or that has an activity against a biological pathway containing a gene required for cellular proliferation or the biological pathway in which a proliferation, or that inhibits cellular proliferation of an organism acts; (9) manufacturing an antibiotic; (10) profiling a compound's activity; (11) a culture comprising strains in which the gene conduct is overexpressed or underexpressed; (12) determining the extent convolute each of the strains is present in a compound that inhibits the collication of an organism. The antisense nucleic acids are useful for identifying proteins or screening for homologous nucleic acids required containing the containing the containing the containing containing the containing containing the containing containing the containing 
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06-SEP-2001; 2001US-00948993.
25-OCT-2001; 2001US-0342923P.
08-FEB-2002; 2002US-00072851.
06-MAR-2002; 2002US-0362699P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 25; SEQ ID NO 55837; 1766pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense nucleic acids, useful for identifying proteins or screening for homologous nucleic acids required for cellular proliferation to isolate candidate molecules for rational drug discovery programs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense; prokaryotic essential gene; cell proliferation; drug design
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dentifying proteins or screening for homologous nucleic a cellular proliferation to isolate candidate molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention relates to an isolated nucleic acid comprising any one of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ELITRA PHARM INC.
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Trawick
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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Carr (
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100.0%; Pred. No.
Live 0; Mismatc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ξ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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Forsyth
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Xu HH;
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RESULT 40
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 25-FEB 1999
05-MAR 1999
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25-MAR 1999
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06-APR 1999
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06-MAY 1999
11-MAY 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 drug discovery programs, or for screening homologous nucleic acids required for proliferation in cells other than S. aureus, S. typhimurium, K. pneumoniae or P. aeruginosa. The present sequence is encoded by one of the target prokaryotic essential genes. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein identification; signal transduction pathway; metabolic pathway; hybridisation assay; genetic mapping; gene expression control; promoter; termination sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAG49871;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAG49871 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Arabidopsis thaliana protein fragment SEQ ID NO: 63135.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Arabidopsis thaliana.
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19 YASNLH
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5; Conserv
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ilarity 83.3%;
Conservative
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99US-0121825P
99US-0123548P
99US-0125788P
99US-0126264P
99US-0126785P
99US-012871462P
99US-01287149
99US-01300479P
99US-0130510P
99US-0130449P
99US-0130449P
99US-0132448P
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99US-013533P
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Pred. No. 4e+02;
1; Mismatches
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27-MAY-1999;
28-MAY-1999;
01-JUN-1999;
03-JUN-1999;
07-JUN-1999;
08-JUN-1999;
10-JUN-1999;
 16-JUN-1999
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99US-014933P
99US-014931P
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PR 03-AUG-1999; 99US-0147038P.
PR 04-AUG-1999; 99US-0147702P.
PR 04-AUG-1999; 99US-0147702P.
PR 05-AUG-1999; 99US-01477303P.
PR 05-AUG-1999; 99US-01477303P.
PR 06-AUG-1999; 99US-01477303P.
PR 06-AUG-1999; 99US-01477303P.
PR 10-AUG-1999; 99US-0148311P.
PR 11-AUG-1999; 99US-0149368P.
PR 12-AUG-1999; 99US-0149368P.
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PR 22-AUG-1999; 99US-0149300P.
PR 22-AUG-1999; 99US-0151066P.
PR 22-AUG-1999; 99US-015106
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PR 28-OCT-1999; 99US-0161993P.
PR 29-OCT-1999; 99US-0162142P.

Query Match
Best Local Similarity 71.4%; Pred. No. 4.4e+02;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Oy 1 YSSNLHS 7

Db 100 FSTNLHS 106
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Search completed: December 17, 2004, 18:29:05 Job time : 23.2753 secs

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Title:
Perfect score:
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2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

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4: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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RESULT 2
US-09-225-322B-19
; Sequence 19, Application US/09225322B
; Patent No. 6437098
; GENERAL INFORMATION:
; APPLICANT: SHITARA, KENYA
; APPLICANT: HANAI, NOBUO

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TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/225,322B
CURRENT FILLING DATE: 1999-01-05
PRIOR FILLING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1994-08-17
PRIOR APPLICATION NUMBER: US 08/297,674
PRIOR APPLICATION NUMBER: US 07/947,674
PRIOR APPLICATION NUMBER: US 07/947,674
PRIOR APPLICATION NUMBER: US 07/947,674
PRIOR APPLICATION NUMBER: US 08/292,178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Description of Artificial Sequence: light chain
; OTHER INFORMATION: variable region
US-09-225-3228-19
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US-09-764-304-10
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 128
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SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 19
LENGTH: 128
TYPE: PRT
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Best Local
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APPLICANT: HANAI, NOBMO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 10, Application US/09764304
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CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER FILING DATE: 1999-01-05
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER FILING DATE: 1995-05-31
                                                                                                                                                                                                                                                                                 EARLIER APPLICATION NUMBER: US07/947,674
EARLIER FILING DATE: 1992-09-17
EARLIER APPLICATION NUMBER: JP 3-238375
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EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER FILING DATE: 1994-08-17
                                                                                                           LENGTH: 12
TYPE: PRT
                                                                                                                                                                                                                                                          EARLIER FILING DATE: 1991-09-18
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                          OTHER INFORMATION: cDNA KM-641
                                                     ORGANISM: Artificial Sequence FEATURE:
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MIYAJI, HJ
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HIROMASA
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100.0%; Pred. No. 5.2
cive 0; Mismatches
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Matches 7
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Patent No. 5
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GENERAL INFORMATION:
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Best Local
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EARLIER FILING DATE: 1992-09-17
EARLIER APPLICATION NUMBER: JP 3-238375
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CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER FILING DATE: 1999-01-05
                                                                                                                                                                                                       GENERAL INFORMATION:
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EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER FILING DATE: 1994-08-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: SHITARA, KEN
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAI
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NUMBER OF SEQ ID NOS: 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 128
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                          APPLICANT:
                              CORRESPONDENCE ADDRESS
                                             APPLICANT: COLMAN, PETER M.
TITLE OF INVENTION: TARGET BINDING POLYPEPTIDE
NUMBER OF SEQUENCES: 25
                                                                                                                          APPLICANT:
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                 ADDRESSEE:
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3: Foley & Lardner
                                                                                           MALBY,
POWER,
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                                                                                                                                     IRVING, Robert A.
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                                                                                           Barbara E.
                                                                                                          Robyn L.
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; SOFTWARE: PatentIn version 2.0
; SEQ ID NO 24
; LENGTH: 474
; TYPE: PRT
; ORGANISM: Cytophaga lytica
US-09-823-823-24
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 273 Amir
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  Best Local Similarity
                    Query Match
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/AU93/00491
FILING DATE: 24-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA: 25-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                     APPLICANT: Suzuki, Makoto
APPLICANT: Hamoda, Tohru
TITLE OF INVENTION: METHOD FOR IDENTIFICATION AND DETECTION OF MICROORGANISMS USING
TITLE OF INVENTION: GENE AS AN INDICATOR
FILE REFERENCE: 12817-004001
CURRENT APPLICATION NUMBER: US/09/823,823
CURRENT FILING DATE: 2001-03-30
RIOR APPLICATION NUMBER: US 09/208,688
PRIOR APPLICATION NUMBER: US 09/208,688
PRIOR APPLICATION NUMBER: JP 97/343316
PRIOR APPLICATION NUMBER: JP 97/343316
PRIOR FILING DATE: 1997-12-12
RUMBER OF SEQ ID NOS: 80
COCTUBER OF SEQ ID NOS: 80
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            PPLICANT: Yamamoto, Satoshi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 16:
TELECOMOUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEPHAX: (202)672-5399
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 30-MAY-19
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Washington STATE: D.C.
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Nakamura, Shoko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/09823823
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85.7%;
89.2%;
85.7%;
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Pred. No. 47;
Score 33; DB 4;
Pred. No. 1.3e+02;
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                    Length 474;
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Query Match
Best Local Similarity
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Patent No. 6455265
GENERAL INFORMATION:
APPLICANT: SERRES, PIERRE-FRANCOIS
APPLICANT: SERRES, PIERRE-FRANCOIS
TITLE OF INVENTION: METHOD FOR OBTAINING VACCINES FOR PREVENTING THE
TITLE OF INVENTION: PATHOGENIC EFFECTS RELATED TO A RETROVIRAL INFECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 4,
Patent No.
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CURRENT FILING DATE: 2000-05-15
PRIOR APPLICATION NUMBER: PCT/FR98/02447
PRIOR PILING DATE: 1998-11-17
PRIOR APPLICATION NUMBER: FR/97/14387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: LashDroux,
APPLICANT: Giovanni, James
TITLE OF INVENTION: Endo-1,4-beta-
TITLE OF INVENTION: Use in Plants
TITLE OF SKOUENCES: 10
                                                                                                                                                                                                                                                                                                                          NUMBER OF SOUTH TOWNSEND ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew
ADDRESSEE: Market Plaza, Steuart Street Tower
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
                CLASSIFICATION: 536
PRIOR APPLICATION UNBER: US 08/271,883
FILING DATE: 07-JUL-1994
PRIOR APPLICATION NDATA:
APPLICATION NUMBER: US 07/687,466
                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                      STREET: One Market I
                                                                                                                         APPLICATION NUMBER: US/0 FILING DATE: 04-MAY-1995
                                                                                                                                                                                                                                                                                        COUNTRY:
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MBER: US 07/687,466
18-APR-1991
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71.4%;
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Endo-1,4-beta-Glucanase Genes and Their
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Pred. No.
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US-09-816-028A-17; Sequence 17, Application US/09816028A; Patent No. 6699705; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                           RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Campylobacter jejuni 
US-09-495-406-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; MOLECULE TYPE: protein US-08-434-702-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: Wakarchuk, Warren W.
APPLICANT: Wakarchuk, Warren W.
APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
TITLE OF INVENTION: Cangylobacter Glycosyltransferases for Biosynthesis of
TITLE REFERENCE: 01963-00011018
CURRENT APPLICATION NUMBER: US/09/495,406
CURRENT FILING DATE: 2000-01-31
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR FILING DATE: 1999-02-01
NUMBER OF SEQ ID NOS: 35
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 13
LENGTH: 347
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 34,774
REPERENCE/DOCKET NUMBER: 022
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
TELEPAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 4:
                     TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis TITLE OF INVENTION: Gangliosides and Ganglioside Mimics FILE REFERENCE: 019633-000111US CURRENT APPLICATION NUMBER: US/09/816,028A CURRENT FILING DATE: 2001-03-21
                                                                                                                       APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 20-APR-
                                                                                                                                                                                                                                                                                                                                                                                                                                            / Match 83.8%;
Local Similarity 71.4%;
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    APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                        268 FSSNIHS 274
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2001-03-21
ER: US 60/118,213
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Pred. No. 2
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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2e+02;
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APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
APPLICANT: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Bio
TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
FILE REPERENCE: 019633-000111US
CURRENT APPLICATION NUMBER: US/10/303,162
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: PatentIn Ver: 2.1
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Best Local Similarity
Thehes 5; Conserve
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US-09-248-796A-22868
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Sequence 22868, Application US/09248796A

Patent No. 6747137

GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANE
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 17
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Patent No. 6723545
                                                                                                                                                                                                                                                                                                                         Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Campylobacter jejuni
FEATURE:
OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNAc)
OTHER INFORMATION: transferase from C. jejuni strain OH4384
OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis
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TYPE: PRT
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ORGANISM: Campylobacter jejuni
FEATURE:
OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNAc)
OTHER INFORMATION: transferase from C. jejuni strain OH4384 (
OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis
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71.4%;
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Pred. No. 2.2e+02;
2; Mismatches 0
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Pred. No.
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RESULT 14
US-09-270-767-45029
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; LENGTH: 67
; TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-22868
                                                                  ; ORGANISM: Drosophila melanogaster US-09-270-767-45029
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US-09-270-767-41649
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Patent No. 6703491
GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
                                                                                                                     APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 45029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 41649
LENGTH: 172
                                                                                                                                                                                                                                                                                            Sequence 45029, Appearent No. 6703491
Best Local Similarity 85. 
Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Drosophila melanogaster FEATURE:
                                                                                                   LENGTH: 238
TYPE: PRT
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                81.1%;
85.7%;
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83.3%;
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83.3%;
Score 30; DB Pred. No. 2.3e 0; Mismatches
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Pred. No. 1.7e+02;
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Pred. No. 65;
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2.3e+02;
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FILE REFERENCE: 15966-542
CURRENT APPLICATION NUMBER: US/09/538,092
CURRENT FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: 60/127,352
PRIOR PILING DATE: 1999-04-01
PRIOR APPLICATION NUMBER: 60/178,965
PRIOR FILING DATE: 2000-02-01
NUMBER OF SEQ ID NOS: 1387
SOFTWARE: CURAPATS SQFTWARE: CURAPATS SQFTWARE SQFTWA
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US-09-252-991A-28155
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                                                                                                                                           Matches
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Patent No. 6753314
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APPLICANT: Mansfield, Traci A.
TITLE OF INVENTION: Protein-Protein Complexes and Method of Using Same
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACII
TITLE OF INVENTION: ABBUGINOSA FOR DIAGNOSTICS
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                                                                                                                                                                                                                                                                                                    LOCATION: (0)...(0)
OTHER INFORMATION: Polypeptide Accession Number Q16600
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ORGANISM: Homo sapiens
FEATURE:
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ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature LOCATION: (0)...(0)
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nes 5; Conserv
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                                                                     2 SSNLHS 7
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NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
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83.3%;
                                                                                                                                                                           81.1%; Score 30;
100.0%; Pred. No.
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Pred. No. 4.1e+02
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RESULT 17

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US-08-630-915A-30

; Sequence 30, Application US/08630915A

; Patent No. 6309820
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US-09-328-352-5898
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Patent No. 6559294
Patent No. 6
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APPLICANT: Gary L. Breton et al.
APPLICANT: GARY L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 8252
SEQ ID NO 5898
LENGTH: 741
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 5898, Application US/09328352 Patent No. 6562958
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Best Local
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                                                        GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.
APPLICANT: HOFFMAN, No. 6309820h
APPLICANT: KAY, Brian K.
APPLICANT: KAY, Erian K.
APPLICANT: MCCONNELL, Stephen J.
APPLICANT: MCCONNELL, Stephen J.
TITLE OF INVENTION: DOMAIN OF INTEREST AND METHODS OF IDENTIFYING.
TITLE OF INVENTION: USING SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: SITE
LOCATION: 1...609
OTHER INFORMATION: Xaa=unknown or other
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les 6; Conserv
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                                     OF SEQUENCES:
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ADDRESS:
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100.0%; Pr/
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83.3%;
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Pred. No. 6.1e+02;
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Pred. No. 7.4e+02;
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GENERAL INFORMATION:
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Best Local Similarity
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CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
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ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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TELECOMMUNICATION INFORMATION:
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STREET: 11
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OPERATING SYSTEM:
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STATE: New Yor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/630,915A FILING DATE: 03-APR-1996
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                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA: Release #1.0,
APPLICATION NUMBER: US/09/879,957
FILING DATE: 13-Jun-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                        COUNTRY: USA
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                       FOWLKES, Dana M.

MCCONNELL, Scephen J.

TITLE OF INVENTION: POLVEEPTIDES HAVING A FUNCTIONAL

DOMAIN OF INTEREST AND METHODS OF IDENTIFYING AND
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,915
                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie
                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: SPARKS, Andrew B
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                                                                                                                                                                                                                                STATE: New York
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                                                                                                                                                                                                                                                  CITY: New York
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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55 Avenue of the Americas
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KAY, Brian K.
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Pred. No. 7.9e+02;
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; MOLECULE TYPE: peptide
; SEQUENCE DESCRIPTION: SEQ ID NO: 30:
US-09-879-957-30
                                                    ; STRANDEDNESS:
; TOPOLOGY: 1i;
; MOLECULE TYPE:
US-08-652-558-2
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                            SOPTWARE: Wordberfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/652,55/
FILING DATE: JUNE 6, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/IB94/003
PILING DATE: NOVEMBER 21, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: LIN, A
                                                                                                                                                                     TELEFAX: 617-345-9111
INFORMATION FOR SEQ ID NO: 2:
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                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 107 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                        NAME: YANKWICH, LEON R.
REGISTRATION NUMBER: 30,237
REFERENCE/DOCKET NUMBER: 95
FELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                      TYPE: amino acid
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ATTORNEY/AGENT INFORMATION:
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TYPE: amino acid
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REFERENCE/DOCKET NUMBER: 1101-174
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (212) 869-8864/9741
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75 STATE STREET, 23RD FLOOR
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VENTION: HUMANIZED ANTIBODIES AND USES
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Conservative
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 78.4%;
71.4%;
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 Score 29; DB 2;
Pred. No. 1.6e+02;
                  Length 107;
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US-09-254-189-1
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MEDIUM TYPE: 3.5 inch diskette, 1.4 Mb storage

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WordPerfect 6.1

CURRENT, APPLICATION DATA:
APPLICATION NUMBER: US/08/652,558

FILING DATE: JUNE 6, 1996

PRIOR APPLICATION NUMBER: PCT/1894/00387

FILING DATE: NOVEMBER 21, 1994

ATTORNEY/AGENT INFORMATION:

REGISTRATION NUMBER: 30,237

ARBISTRATION NUMBER: 30,237
APPLICANT: Lundquist, Tomas
TITLE OF INVENTION: Sequence Listing
Patent No. 6150792
PILE REFERENCE: 3526/00000
CURRENT PILING DATE: 1999-03-02
EARLIER APPLICATION NUMBER: 00/000,000
EARLIER FILING DATE: 1998-11-24
                                                                                                                                    Sequence 1, Application US/09254189 Patent No. 6150792 GENERAL INFORMATION:
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Best Local Similarity
Matches 5; Conserv
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TELEPAX: 617-345-9111
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
LENGTH: 107 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 95
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: LIN, AUGUSTINE YEE-THARN
TITLE OF INVENTION: HUMANIZED ANTIBODIES AND USES
TITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: BOSTON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                        50 YTSSLHS 56
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71.4%;
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Pred. No. 1.6e+02;
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Query Match
Best Local Similarity
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; OTHER INFORMATION: monoclonal antibody
US-09-254-189-1
                                                                              ; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-21191
                                                                                                                                                                                                                                                                                                                                                                                                US-09-252-991A-21191
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APPLICANT: Homburger et al.
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins
FILE REFERENCE: File Reference: 7326-094
CURRENT EPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: PALENTIN Ver. 2.0
                                                                                                             PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 21191
LENGTH: 329
TYPE: PRT
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LENGTH: 219
TYPE: PRT
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Best Local (
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Best Local Similarity
                                                                                                                                                                                                                                           APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
OTHER INFORMATION: Xaa means
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71.4%;
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57.1%;
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71.4%;
             ; Score 29; DB;
; Pred. No. 5e+0
1; Mismatches
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; Pred. No. 1.6e
2; Mismatches
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Pred. No. 3.3e+02;
Pred. No. 3.3e+02;
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. 1.6e+02;
                              5e+02;
                                                DB 4;
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                                                Length 329;
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US-09-710-279-1386
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APPLICANT: LYNIN DOUGETTE-Stamm et al
APPLICANT: LYNIN DOUGETTE-STAMM et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
TITLE REFERENCE: GTC-007
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1997-11-08
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5674

SEQ ID NO 4411
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PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 1386
LENGTH: 496
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Patent No. 670349
                                                                                      Query Match
Best Local Similarity
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TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PUJ480US
CURRENT APPLICATION NUMBER: US/09/710,279
CURRENT FILING DATE: 2000-11-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Artificial Sequence
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                                                                                                                                                                                                     LENGTH: 497
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275 YDSNLH 280
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                                    1 YSSNLH
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83.3%;
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83.3%;
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Pred. No. 7.6e+02
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Pred. No.
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                                                                          Mismatches
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7.6e+02;
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RESULT 28 US-08-522-269B-3

Sequence 3, Application US/08522269B Patent No. 5919690

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RESULT 29
US-09-294-923-3
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                                                                                                                                                                                                                                                                                                                                                                                                         Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 212 867 0298
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 677 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Knap, Inge Helmer
APPLICANT: Hjort, Carsten M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
            CITY: New York

CITY: New York

COUNTRY: U.S.A.

ZIP: 10174-6401

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)

CURRENT APPLICATION DATA:
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NAMES Gregg, Valeta A.
REGISTRATION NUMBER: 37,125
REFERENCE/DOCKET NUMBER: 39:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 2
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                                                                                                                                                                                                                                                                   APPLICANT: Halkier, Torben
APPLICANT: Kofod, Lene Venke
IITLE OF INVENTION: An Alpha-Galactosidase Enzyme
NUMBER OF SEQUENCES: 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PPLICANT: Halkier, Torben
PPLICANT: Kofod, Lene Venke
ITLE OF INVENTION: An Alpha-Galactosidase Enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        78.4%;
Local Similarity 71.4%;
es 5; Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/522,269B FILING DATE: 12-SEP-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A. ZIP: 10174-6401
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APPLICATION NUMBER:
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                                                                                                                                                                                                          405 Lexington Avenue, Suite 6400
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Hjort, Carsten M.
Halkier, Torben
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linear
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US/09/294,923
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Pred. No. 1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1e+03;
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GENERAL INFORMATION:
APPLICANT: Allen, Steve
APPLICANT: Allen, Steve
APPLICANT: Kinney, Tony
APPLICANT: Kinney, Tony
APPLICANT: Tingey, Scott
TITLE OF INVENTION: Plant Sugar Transport Proteins
FILE REFERENCE: BB-1163
CURRENT APPLICATION NUMBER: US/09/291,922
CURRENT FILING DATE: 1999-04-14
EARLIER APPLICATION NUMBER: 60/083,044
EARLIER APPLICATION NUMBER: 60/083,044
EARLIER FILING DATE: APPII 24, 1998
NUMBER OF SEQ ID NOS: 30
SOFTWARE: Microsoft Office 97
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          밁
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/09291922
Patent No. 6383776
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -09-291-922-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 212 867 0298
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 677 amino acids
                                                                                                                                                                                                                                                                                                            LENGTH: 747
TYPE: PRT
ORGANISM: Zea mays
FEATURE:
                                                                                                                                                                                                                                                                               NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                  NAME/KEY:
LOCATION:
                                                                                                                                                                                                     VAME/KEY:
                                                                                                                                                                                                                                                                  EATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Gregg, Valeta A.
REGISTRATION NUMBER: 37,125
                AME/KEY:
                                EATURE
                                                              AME/KBY:
                                                                             EATURE
                                                                                                            IAME/KEY
                                                                                                                                          OCATION:
                                                                                                                                                                                        OCATION:
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                                                                                             OCATION:
                                                                                                                                                                                                                      EATURE
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                                              OCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 12-SE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         212 YSSHLHN 218
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(129)
UNSURE
(220)
                                                                                          UNSURE
(207)
                                                                                                                                       UNSURE
(178)
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(144)
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(133)..(134)
                                             (218)
                                                           UNSURE
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71.4%;
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; LOCATION: (236)
US-09-291-922-2
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US-08-137-117D-118
                                                                                                                       ; TOPOLOGY: ]
US-08-137-117D-118
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Best Local Similarity 71.4%;
Matches 5; Conservative
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                                                                     Query Match
Best Local Similarity
                                                   Matches
                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION UNMBER: WO PCI
FILING DATE: 24-APR-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 20007-5109
ZOMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
TOTAL TAMERON: 100/00/127 1170
                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                         NAME: WEGNER, Harold C.
REGISTRATION NUMBER: 25,258
REFERENCE/DOCKET NUMBER: 53
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: JP 4-32084
FILING DATE: 19-FEB-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 3-95476
                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: JP 3-95476 FILING DATE: 25-APR-1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: RESHAPED HUMAN ANTIBODY TO HUMAN TITLE OF INVENTION: INTERLEUKIN-6 RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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TELEFAX: 904136
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                     LENGTH: 7 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/137,117D FILING DATE: 20-DEC-1993
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CITY: Washington
STATE: D.C.
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3000 K Street, N.W., Suite 500
                                                     Conservative
                                                                                                                                                                                                                                                           (202) 672-5399
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BENDIG, Mary
JONES, Steven
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                                                                                                                                       linear
                                                                     71.4%;
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                                                 Score 28; DB 1; 1
Pred. No. 3.8e+05;
1; Mismatches 1.
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Pred. No. 1.2e+03;
0; Mismatches 2;
                                                                                   Length 7;
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                                                     1; Indels
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Query Match
Best Local Similarity
Tatches 5; Conserve
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Patent No. 5811248
GENERAL INFORMATION:
APPLICANT: Charle
                                                                                                                                                                                 Patent No. 5817790 GENERAL INFORMATION:
                                                                                                                                                                                                  Sequence 118, Application US/08436717 Patent No. 5817790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY AGENT INFORMATION:
NAME: Albert P. Halluin
REGISTRATION NUMBER: 25,227
REFERENCE/DOCKET NUMBER: 7606
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 854-3660
TELEFAX: (415) 854-3694
                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: D
                                                              APPLICANT: BENDIG, Mary
APPLICANT: JONES, Steven
APPLICANT: SALDANHA, JOSE
TITLE OF INVENTION: RESHAPED HUMAN ANTIBODY TO HUMAN
TITLE OF INVENTION: INTERLEUKIN-6 RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     HYPOTHETICAL:
ANTI-SENSE: N
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CLASSIFICATION: 43:
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ZIP: 10036
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STATE: New York
 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
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                  ADDRESSEE:
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E: Foley & Lardner
3000 K Street, N.W., Suite 500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7 amino acids
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                                                                                                                                                  TSUCHIYA, Masayuki
SATO, Koh
                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JUN-1995
N: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       unknown
                                                                                                                                                                                                                                                                                                                                                                                     75.7%;
71.4%;
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                                                   158
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                                                                                                                                                                                                                                                                                                                                                                                  Score 28; DB 2;
Pred. No. 3.8e+05;
                                                                                                                                                                                                                                                                                                                                                                      1; Mismatches
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RESULT 34
US-08-053-451B-78
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APPLICANT: Chen, Fre
APPLICANT: Ditlow, C
APPLICANT: Calenoff,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 78, Application US/08053451B Patent No. 5955584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 536

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/137,117

FILING DATE: 20-DEC-1993
APPLICATION NUMBER: WO PCT/JP92/005-
FILING DATE: 24-APR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 4-32084
FILING DATE: 19-FEB-1992
PRIOR APPLICATION DATA:
APPLICATION UMBER: JP 3-95476
FILING DATE: 25-APR-1991
ATTORNEY/AGENT INFORMATION:
ANDEE: MECNIE 1902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 904136
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
               COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 53
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (202)672-5399
                                                                                                                                              STATE: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              / Match 75.7%;
Local Similarity 71.4%;
                                                                                                                                                             CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: WEGNER, HATOLO REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                  STREET:
APPLICATION NUMBER:
                                                                                                                           10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 YSSNLHS 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7 amino acids
                                                                                                                                                                                                  1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                      Calenoff, Emanue.
                                                                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     single
                                                                                                                                                                                                                                                                                                                                 Charles C.
                                                                                                                                                                                                                                                                         ATHEROSCLEROTIC PLAQUE SPECIFIC ANTIGENS, ANTIBODIES THERETO, AN
                                                                                                                                                                                                                                                           176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO PCT/JP92/00544
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US/08/053,451B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 28; DB 2;
Pred. No. 3.8e+05;
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    Indels

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 7;
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                                            TELEFAX: (703) 205-8000
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
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Best Local 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: MURPHY JR, GERALD M
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 1110-160
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/649,100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: ANTI-FAS LIGAND ANTIBODY AND ASSAY TITLE OF INVENTION: METHOD USING THE ANTI-FAS LIGAND ANTIBODY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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TOPOLOGY: li
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STREET: PO
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REFERENCE/DOCKET NUMBER: 7606-033-999
                            TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                              CITY: FALLS CHURCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: 415-854-3694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 26-APR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13,
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PO BOX 747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VASQUEZ, MAXIMILIANO
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           sss: single
linear
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MATUSUE, TOMOKAZU
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                                                                                                                                                                                                                                                                                                                                                                                                                           STEWART, KOLASCH AND BIRCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                           41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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US-09-563-222C-39; Sequence 39, Application US/09563222C; Patent No. 6696620; Patent INFORMATION:
; APPLICANT: EPICYTE PHARMACEUTICALS, INC.; APPLICANT: HIATT, ANDREW C.; APPLICANT: HEIN, MICH B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 36
US-08-649-100-29
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Best Local Similarity
Watches 5; Conserv
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                                                                                                                                                                                                                                                       RESULT 37
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ATTORNEY/AGENT INFORMATION:
NAME: MURPHY JR, GERALD M
REGISTRATION NUMBER: 28,977
REFERENCE/DOCKET NUMBER: 1110
TELECOMMUNICATION INFORMATION:
TELEBAX: (703) 205-8000
TELEFAX: (703) 205-8050
INFORMATION FOR SEQ ID NO: 29:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/649,100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: VASQUEZ, MAXIMILIANO
TITLE OF INVENTION: ANTI-FAS LIGAND ANTIBODY AND ASSAY
TITLE OF INVENTION: METHOD USING THE ANTI-FAS LIGAND ANTIBODY
NUMBER OF SEQUENCES: 41
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
CITY: F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 22040-0747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: VA
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YTSRLHS 7
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MATUSUE, TOMOKAZU
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71.4%;
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71.4%; Pred. No. 3.8e+05;
tive 1; Mismatches 1
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Pred. No. 3.8e+05;
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                                                                               US-08-525-539A-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: IMMUNOGLOBULIN BINDING PROTEIN ARRAYS IN PLANT CELLS
FILE REFERENCE: 068994-0501
CURRENT APPLICATION NUMBER: US/09/563,222C
CURRENT FILING DATE: 2000-05-02
PRIOR APPLICATION NUMBER: PCT/US01/14349
PRIOR APPLICATION NUMBER: PCT/US01/14349
PRIOR APPLICATION NUMBER: 09/563,222
PRIOR APPLICATION NUMBER: 09/563,222
PRIOR FILING DATE: 2000-05-02
NUMBER OF SEQ ID NOS: 182
SEQ ID NO 39
                                                                                                                                        TELEX: 706141
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 14, Application US/08525539A Patent No. 6309636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 75.7%;
Best Local Similarity 71.4%;
Matches 5; Conservative
Matches
                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:

APPLICANT: DO COUTO, FERNANDO J.R.

APPLICANT: CERIANI, ROBERTO L.

APPLICANT: PETERSON, JERRY A.

APPLICANT: PETERSON, JERRY A.

TITLE OF INVENTION: RECOMBINANT PEPTIDES DERIVED FROM THE

TITLE OF INVENTION: MC3 ANTI-BA46 ANTIBODY, METHODS OF USE THEREOF, AND

TITLE OF INVENTION: METHODS OF HUMANIZING ANTIBODY PEPTIDES
                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: DYLAN, TYLER
REGISTRATION NUMBER: 37,612
REFERENCE/DOCKET NUMBER: 2763
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 7
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: PAt-entin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/525,539A
FILING DATE: 14.5EP-1995
CLASSIFICATION: 424
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                  LENGTH: 32 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 755 Page Mill Road CITY: Palo Alto
                                                                                                                                                                                                                      TELEPHONE: (415) 494-0792
                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: MORRISON & FOERSTER
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  Conservative
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71.4%;
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                    Score 28;
Pred. No.
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  Mismatches
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                                      Length 32;
  0;
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    Gaps
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Sequence 19786, Application US/09248796A

Patent No. 6747137

GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
ITILE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
ITILE OF INVENTION: NUCLEIC ACID AND THERAPEUTICS

FILE REFERENCE: 107196, 132

CURRENT APPLICATION NUMBER: US/09/248,796A

CURRENT FILING DATE: 1999-02-12

PRIOR APPLICATION NUMBER: US 60/074,725

PRIOR APPLICATION NUMBER: US 60/074,725

PRIOR APPLICATION NUMBER: US 60/096,409

PRIOR FILING DATE: 1998-08-13

NUMBER OF SEQ ID NOS: 28208

SEQ ID NO 19786

LENGTH: 99

TYPE: PRT
ORGANISM: Candida albicans

US-09-248-796A-19786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 5271, Application US/09134000C

Patent No. 6617156

GENERAL INFORMATION:

APPLICANT: Lynn Doucette-Stamm et al

APPLICANT: Lynn Doucette-Stamm et al

ITITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO

ITITLE OF INVENTION: ENTEROCOCCUS FASCALIS FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 032796-032

CURRENT APPLICATION NUMBER: US/09/134,000C

CURRENT FILING DATE: 1998-08-13

PRIOR APPLICATION NUMBER: US 60/055,778

PRIOR APPLICATION NUMBER: US 60/055,778

PRIOR FILING DATE: 1997-08-15

NUMBER OF SEQ ID NOS: 6812

SOFTWARE: Patentin version 3.1

LENGTH: 98

TYPE: PRT
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Search completed: December 17, 2004, 18:32:09 Job time: 4.24719 secs
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US-09-248-796A-19786
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US-09-134-000C-5271
                                                                                                                                                                                                  Query Match 75.7%; Score 28; DB 4; Length 99; Best Local Similarity 83.3%; Pred. No. 2.3e+02; Matches 5; Conservative 1; Mismatches 0; Indels
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Minimum
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185.159 Million cell updates/sec
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37
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/cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
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/cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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33	33	33	33	33	34	34	37	37	37	37	37	37	Score
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474	474	306	303	91	2207	107	128	128	128	128	128	128	Query Match Length DB
9	9	15	15	15	16	17	14	14	14	14	9	9	BB
US-09-823-823-24	US-09-823-829-24	US-10-425-114-49629	US-10-424-599-208863	US-10-424-599-201107	US-10-437-963-189767	US-10-741-657A-16	US-10-166-626-19	US-10-166-626-10	US-10-265-713-19	US-10-265-713-10	US-09-764-304-19	US-09-764-304-10	ID
Sequence 24, Appl	Sequence 24, Appl	Sequence 49629, A	Sequence 208863,	Sequence 201107,	Sequence 189767,	Sequence 16, Appl	Sequence 19, Appl	70,	Sequence 19, Appl		Sequence 19, Appl	Sequence 10, Appl	Description

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14
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0-424-599-16530	-10-425-115-18936	0-425-115-19318	-10-425-115-22353	-10-424-599-26424	-10-425-115-24254	0-424-599-23154	-10-425-1	-10-424-599-28557	-10-425-114-6436	-10-425-114-4902	-10-739-930-762	-10-425-115-320	-10-847-983-	0-821-604-	-10-821-573-1	-10-846-	0-845-412-	-10-845-408-	-10-820-536-1	-10-303-162-	-10-303-134-	-10-303-128-1	0-303-118-	0-303-161-1	09-816-028A-	0-282-122A-5338	-10-425-115-31	-09-970-36	-10-424-599-19005	-10-425-115-227	0-437-963-19700
16530	18936	e 193	22353	e 2642	24254	23154	Sequence 192258,	28557	"	e 49028,	e 7620	e 320	e 17,	e 17,	e 17,	Sequence 17, Appl	17,	17,	,7	17,	17,	equence 17,	e 17,	e 17,	17,	e 533	æ	Φ	190057,	e 22777	19

## ALIGNMENTS

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RESULT 1
US-09-764-304-10
Sequence 10, Application US/09764304
Patent No. US20020026036A1
GENERAL INFORMATION:
APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HANAI, NOBUO
APPLICANT: HANAI, NOBUO
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUWANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 199-01-05
EARLIER FILING DATE: 1995-05-31
EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER FILING DATE: 1995-03-21
EARLIER FILING DATE: 1992-09-17
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER PILING DATE: 1991-09-18

SOPTMARE: PARTICATION NUMBER: US 08/408,133
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER PILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER PILING DATE: 1995-03-21
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EARLIER PILING DATE: 1995-03-21
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EARLIER PILING DATE: 1995-03-21
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EARLIER PILING DATE: 1995-03-21
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Query Match

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Length 128;

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CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER FILING DATE: 1999-01-05
EARLIER FILING DATE: 1999-01-05
EARLIER FILING DATE: 1995-05-31
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER FILING DATE: 1995-03-21
EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER APPLICATION NUMBER: US07/947,674
EARLIER FILING DATE: 1994-08-17
EARLIER APPLICATION NUMBER: US07/947,674
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                                                                                                                                                                                                      Sequence 10, Application US/10265713

Publication No. US2003095964A1

GENERAL INFORMATION:
APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 19
LENGTH: 128
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/265,713
CURRENT FILING DATE: 2002-10-08
PRIOR APPLICATION NUMBER: US/09/225,322
PRIOR FILING DATE: 1999-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
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APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
                                                                                                                                                                                            FILE REFERENCE: 249-101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
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100.0%; Pred. No.
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PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US07/947,674
PRIOR FILING DATE: 1992-09-17
PRIOR APPLICATION NUMBER: UF 3-238375
PRIOR FILING DATE: 1991-09-18
NUMBER OF SEO ID NOS: 19
SOFTWARE: Patentin Ver. 2.0
SEO ID NO 19
В
                                                                                                                                                        ; OTHER INFORMATION: Description of Artificial Sequence:light chain ; OTHER INFORMATION: variable region US-10-265-713-19
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; OTHER INFORMATION: Description of Artificial Sequence: cDNA KM-641
US-10-265-713-10
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PRIOR FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
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APPLICANT: HANAI, NOSUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAAI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/265,713
CURRENT FILING DATE: 2002-10-08
PRIOR APPLICATION NUMBER: US/09/225,322
PRIOR FILING DATE: 1999-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
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PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1994-08-17
PRIOR APPLICATION NUMBER: US07/947,674
PRIOR FILING DATE: 1992-09-17
PRIOR FILING DATE: 1992-09-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                           LENGTH: 128
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                                     1 YSSNLHS 7
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                                                                            Similarity 7; Conserv
  YSSNLHS
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Pred. No. 21;
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APPLICANT: HANAI, NOBUO
APPLICANT: HANAI, NOBUO
APPLICANT: MIYAGI, HIROMASA
APPLICANT: MIYAGI, PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/10/166,626
CURRENT FILING DATE: 2002-06-12
PRIOR APPLICATION NUMBER: US/09/225,322B
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR FILING DATE: 1995-05-31
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US07/947,674
PRIOR APPLICATION NUMBER: US07/947,674
PRIOR APPLICATION NUMBER: UP30-09-17
PRIOR APPLICATION NUMBER: UP30-09-18
PRIOR FILING DATE: 1991-09-18
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CURRENT FILING DATE: 2002-06-12
PRIOR APPLICATION NUMBER: US/09/225,3228
PRIOR FILING DATE: 1999-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 08/292,179
PRIOR APPLICATION NUMBER: US 08/29375
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Matches
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APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAM
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APPLICANT: HANAI, NOBUO
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ORGANISM: Artificial Sequence
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TILE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
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les 7; Conserv
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SEQ ID NOS: 19
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                                                                                             TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)B CURRENT APPLICATION NUMBER: US/10/437,963 CURRENT FILING DATE: 2003-05-14 NUMBER OF SEQ ID NOS: 204966 SEQ ID NO 189767

LENGTH: 2207
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SOFTWARE: Patentin version 3.2
SEQ ID NO 16
LENGTH: 107
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas
APPLICANT: Kovalic, David
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Matches 6; Conser
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TITLE OF INVENTION: ANTIBODIES AGAINST GPR64 AND USES THEREOF
FILE REFERENCE: 05882.0177.NPUS01
                                                                                                                                                                                                                                 APPLICANT:
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NAME/KEY: unsure
LOCATION: (1)..(2207)
OTHER INFORMATION: unsure at all Xaa locations
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                                                                                 TYPE: PRT
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                                                               ORGANISM: Oryza sativa
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Zhou, Yihua
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Barbazuk, Brad
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Pred. No.
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Pred. No.
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                                                                                                 ; OTHER INFORMATION: Clone ID: PAT_MRT3847_30631C.1.pep
US-10-424-599-208863
                                                                                                                                                                                          APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(5323)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 208863
LENGTH: 303
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LENGTH: 91
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APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53223) B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
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APPLICANT: Kovalic David K
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                                                                                                                                      TYPE: PRT
ORGANISM: Glycine max
FEATURE:
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ORGANISM: Glycine max
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                    Conservative
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85.7%;
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Pred. No. 8
                Score 33; DB 15; Length 303; Pred. No. 2.8e+02; Mismatches 0; Indels
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Pred. No.
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                                                                           Query Match
Best Local S
Matches 6
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Best Local S
Matches 6
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APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven B
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53313) B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 49629
**PROVENUM 106
                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 82
SOFTWARE: Patentin version 2.0
SEQ ID NO 24
LENGTH: 474
TYPE: PRT
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Patent No. US20020146697A1
GENERAL INFORMATION:
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Publication No. US20040034888A1
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                                                                                                                                                                                                                                                                                               APPLICANT: Hamada, Töhru
TITLE OF INVENTION: METHOD FOR IDENTIFICATION AND DETECTION OF MICROORGANISMS
TITLE OF INVENTION: USING GYRASE GENE AS AN INDICATOR
FILE REFERENCE: 12817-004001
CURRENT APPLICATION NUMBER: US/09/823,829
CURRENT FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: US 09/208,688
PRIOR FILING DATE: 1998-12-10
PRIOR APPLICATION NUMBER: US 09/208,688
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ORGANISM: Glycine max
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164 YSENLHS 170
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                                                                             Similarity 6; Conserv
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Suzuki, Makoto
Kasai, Hiroaki
                                     YSSNLHS 7
                                                                           Conservative
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85.7%;
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85.7%;
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Pred. No.
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                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 197006
LENGTH: 503
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SOFTWARE: PatentIn version 2.0
SEQ ID NO 24
                                                                                                               Query Match
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                                                                                              Best Local Similarity
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Patent No. US20020171092A1
                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(53221)B CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -09-823-823-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/208,688
PRIOR FILING DATE: 1998-12-10
PRIOR APPLICATION NUMBER: JP 97/343316
PRIOR FILING DATE: 1997-12-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: METHOD FOR IDENTIFICATION AND DETECTION OF MICROORGANISMS USING TITLE OF INVENTION: GENE AS AN INDICATOR FILE REFERENCE: 12817-004001
CURRENT APPLICATION NUMBER: US/09/823,823
CURRENT FILING DATE: 2001-03-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
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ublication No. US20040123343A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-437-963-197006
                                                                                                                                                                                               ORGANISM: Oryza sativa FEATURE:
                                                                                                                                                                        OTHER INFORMATION: Clone ID: PAT_MRT4530_92805C.1.pep
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                                  YSSNLH 6
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YSSNLH 89
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Nakamura, Shoko
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                                                                          Conservative
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85.7%;
                                                                          89.2%; Score 33; DB 16;
100.0%; Pred. No. 4.7e+02;
tive 0; Mismatches 0;
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Pred. No. 4.4e+02;
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RESULT 15

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US-10-424-599-190057
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US-10-424-599-190057
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Best Local S
Matches 6
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CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 190057
LENGTH: 59
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Publication No. US20040031072A1
GENERAL INFORMATION:
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SEQ ID NO 227775
                                                                                                            Sequence 2, Application US/09970367
Publication No. US20030106092A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
                                                       APPLICANT: Davis, APPLICANT: Goell
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                 APPLICANT: Davis, Bric L.
APPLICANT: Goelliner, Melissa
TITLE OF INVENTION: ENDOGLUCANASE GENE PROMOTER UPREGULATED BY NEMATODES
FILE REFERENCE: 5051.587
CURRENT APPLICATION NUMBER: US/09/970,367
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ORGANISM: Zea mays
FEATURE:
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Cao, Yongwei
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Pred. No.
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Pred. No.
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 316851
TUDE: DEFE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TYPE: PRT ; ORGANISM: Nicotiana tabacum US-09-970-367-2
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US-10-282-122A-53384
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US-10-425-115-316851
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NUMBER OF SEQ ID NOS: 2
SOFTWARE: PatentIn vers
SEQ ID NO 2
LENGTH: 500
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                                                                                                                                                                                                                APPLICANT: Wang, I
APPLICANT: Zamudi
APPLICANT: Malond
APPLICANT: Hasell
APPLICANT: Ohlsel
APPLICANT: Zyskii
                                                                                                                                                                                                                                                                                                                          Sequence 53384, Application US/10282122A Publication No. US20040029129A1 GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT: Wall, Daniel
APPLICANT: Trawick, John
APPLICANT: Carr, Grant
APPLICANT: Yamamoto, Robert
APPLICANT: Yamamoto, Robert
APPLICANT: Yamamoto, Robert
APPLICANT: Yau, H.
APPLICANT: You, H.
TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
FILE REFERENCE: ELITRA.034A
CURRENT APPLICATION NUMBER: US/10/282,122A
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Zea mays FEATURE:
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Local Similarity 85.7%;
es 6; Conservaring
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Malone, Cheryl
                                                                                                                                                                                                                    Ohlsen, Kari
Zyskind, Judith
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                                                                           Query Match
Best Local S
Matches 5
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APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis
TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
FILE REFERENCE: 019633-00011US
FILE REFERENCE: 019633-00011US
CURRENT APPLICATION NUMBER: US/09/816,028A
CURRENT FILING DATE: 2001-03-21
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 53384
LENGTH: 277
                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 17
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Best Local (
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PRIOR APPLICATION NUMBER: 60/269,308
PRIOR FILING DATE: 2001-02-16
                                                                                                                                                                        OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNac) OTHER INFORMATION: transferase from C. jejuni strain OH4384 (ORF OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis locu
                                                                                                                                                                                                                               ORGANISM: Campylobacter jejuni
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Clostridium difficile
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FILING DATE: 2000-05-23
APPLICATION NUMBER: 60/
FILING DATE: 2000-05-26
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APPLICATION NUMBER: 60/267,636
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FILING DATE: 2000-11-27
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APPLICATION NUMBER: 60/257,931
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  268 FSSNIHS 274
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                                      1 YSSNLHS 7
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Pred. No.
                                                                                             Score 31; DB 9;
Pred. No. 7.8e+02;
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6.2e+02;
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APPLICANT: National Research Council of Canada
TITLE OF INVENTION: Campylbbacter Glycosyltransferases for Biosynthesis of
TITLE OF INVENTION: Gangylbbacter Glycosyltransferases for Biosynthesis of
TITLE OF INVENTION: Gangylbosides and Ganglioside Mimics
FILE REFERENCE: 019633-000111US
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/09/016,028
PRIOR APPLICATION NUMBER: US/09/016,028
PRIOR APPLICATION NUMBER: US/09/016,213
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 1999-02-01
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER: OF SEQ ID NOS: 49
SOPTWARE: PATENTIN Ver. 2.1
SEQ ID NO 17
LENGTH: 347
TYPE: DET
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US-10-303-118-17
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US-10-303-161-17
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CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR FILING DATE: 1999-02-01
PRIOR FILING DATE: 1909-03-01
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: PATENTIN Ver: 2.1
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APPLICANT: Wakarchuk, War
APPLICANT: National Resea
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APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
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OTHER INFORMATION: transferase from C. jejuni strain OH4384 (ORF 5a
OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis locus)
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TYPE: PRT
ORGANISM: Campylobacter jejuni
FEATURE:
                                                                            OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNAc) OTHER INFORMATION: transferase from C. jejuni strain OH4384 OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis
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71.4%;
  83.8%;
Score 31;
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  DB 14;
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Length 347;
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APPLICANT: Gilbert, Michel
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
APPLICAT: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
FILE REFERENCE: 019633-00011US
CURRENT APPLICATION NUMBER: US/10/303,134
CURRENT FILING DATE: 2002-11-21
                                         NUMBER OF SEQ ID NOS: 49
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 17
LENGTH: 347
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Best Local Similarity 71.4%;
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Matches 5; Conservative
                                                                                                            PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
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CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
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SOFTWARE: Patentin Ver.
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PRIOR FILING DATE: 2000-01-31
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APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNAc) OTHER INFORMATION: transferase from C. jejuni strain OH4384 OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 347
TYPE: PRT
ORGANISM: Campylobacter jejuni
ORGANISM: Campylobacter jejuni
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Sequence 17, Application US/10303162
Publication No. US20030157658A1
GENERAL INFORMATION:
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: Wakarchuk, Warren W.
APPLICANT: Wational Research Council of Canada
ITITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
ITITLE OF INVENTION: Gangliosides and Ganglioside Mimics
FILE REFERENCE: 019633-000111US
CURRENT APPLICATION NUMBER: US/10/303,162
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/9/916,028
PRIOR APPLICATION NUMBER: US/9/918,028
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
PRIOR APPLICATION SUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
VNUMBER OF SEQ ID NOS: 49
SOFTWARE: Patentin Ver. 2.1
LENGTH: 347
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: Wational Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
TITLE OF INVENTION: Cangliosides and Ganglioside Mimics
FILE REFERENCE: 019633-00011US
CURRENT APPLICATION NUMBER: US/10/820,536
CURRENT FILING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR RILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR APPLICATION NUMBER: US/09/816,028
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Matches
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Publication No. US20040203103A1
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNAC) OTHER INFORMATION: transferase from C. jejuni strain OH4384 (ORF 5a OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis locus)
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ORGANISM: Campylobacter jejuni
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Pred. No. 7.8e+02;
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RESULT 28
US-10-845-412-17
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Best Local Similarity
Tatches 5; Conserva
                                                                                            ; Sequence 17, Application US/10845412; Publication No. US20040203113A1; GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 17
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis TITLE OF INVENTION: Gangylobacter Glycosyltransferases for Biosynthesis FILE REFERENCE: 019633-00011185 and Ganglioside Mimics CURRENT APPLICATION NUMBER: US/10/845,408 CURRENT FILING DATE: 2004-05-12 PRIOR APPLICATION NUMBER: US/9/816,028 PRIOR APPLICATION NUMBER: US/9/816,028 PRIOR FILING DATE: 2001-03-21 PRIOR APPLICATION NUMBER: US 60/118,213 PRIOR FILING DATE: 1999-02-01 PRIOR FILING DATE: 1999-02-01 PRIOR FILING DATE: 1999-02-01 PRIOR APPLICATION NUMBER: US 09/495,406 PRIOR APPLICATION STEP 1005: 49 SOFTWARE: PAtentin Ver. 2.1
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases
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APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
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FEATURE:
OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNAc)
OTHER INFORMATION: transferase from C. jejuni strain OH4384 |
OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNAc) OTHER INFORMATION: transferase from C. jejuni strain OH4384 OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 347
TYPE: PRT
ORGANISM: Campylobacter jejuni
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TYPE: PRT
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                                                                                                                                                                                                                                                            268 FSSNIHS 274
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71.4%;
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Pred. No. 7.8e+02;
2; Mismatches 0
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Pred. No. 7.8e+02;
2; Mismatches 0;
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           for Biosynthesis
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US-10-846-219-17
Sequence 17, Application US/10846219
Publication No. US20040219638A1
GENERAL INFORMATION:
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
APPLICANT: National Research Council of Canada
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CURRENT APPLICATION NUMBER: US/10/845,412
CURRENT FILING DATE: 2004-05-12
PRIOR APPLICATION NUMBER: US/10/303,128
PRIOR FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/9/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOPTWARE: PACENTIN Ver. 2.1
                                                                                                                                                                                           US-10-846-219-17
                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis
TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
EILE REFERENCE: 019633-00011US
CURRENT APPLICATION NUMBER: US/10/846,219
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR PILING DATE: 1999-02-01
PRIOR FILING DATE: 2000-01-31
NUMBER OF EGG ID NOS: 49
SOFTWARE: Patentin Ver. 2.11
SEG ID NO 17
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                                                                                                                Query Match
Best Local (
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ORGANISM: Campylobacter jejuni
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Campylobacter jejuni
FEATURE:
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                                                                                                                                                                                                          OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNac)
OTHER INFORMATION: transferase from C. jejuni strain OH4384 (ORF 5a
OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis locus)
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                                                                                                                  Local Similarity
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268 FSSNIHS 274
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                                            YSSNLHS 7
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                                                                                                                83.8%;
                                                                                                                Score 31; DB 17; Length 347; Pred. No. 7.8e+02;
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Pred. No. 7.8e+02;
                                                                                            Mismatches
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APPLICANT: Wakarchuk, Warren W.
APPLICANT: Wakarchuk, Warren W.
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
FILE REFERENCE: 019633-000111US
CURRENT APPLICATION NUMBER: US/10/821,573
CURRENT FILING DATE: 2004-04-08
PRIOR APPLICATION NUMBER: US/9/816,028
PRIOR APPLICATION NUMBER: US/9/816,028
PRIOR APPLICATION NUMBER: US/9/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: PAtentin Ver: 2.1
SEQ ID NO 17
Type: Datentin Ver: 2.1
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PRIOR FILING DATE: 2002-11-21
PRIOR PELICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
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Best Local S
Matches 5
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Publication No. US20040229263A1
GENERAL INFORMATION:
APPLICANT: Gilbert, Michel
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Publication No.
                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 17
LENGTH: 347
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of TITLE OF INVENTION: Gangliosides and Ganglioside Mimics FILE REFERENCE: 019633-000111US CURRENT APPLICATION NUMBER: US/10/821,604
CURRENT FILING DATE: 2004-04-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
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FEATURE: FEATURE: OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNAc) OTHER INFORMATION: transferase from C. jejuni strain OH4384 (OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Campylobacter jejuni
FEATURE:
OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNac)
OTHER INFORMATION: transferase from C. jejuni strain OH4384
OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis
                                                                                                        TYPE: PRT ORGANISM: Campylobacter jejuni
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5; Conserv
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Pred. No.
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ide (LOS) biosynthesis
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7.8e+02;
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APPLICANT: Watarchik, Warren W.
APPLICANT: National Research Council of Canada
ITITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
ITITLE OF INVENTION: Gangliosides and Ganglioside Mimics
FILDE REFERENCE: 0.19633-000111US
CURRENT APPLICATION NUMBER: US/10/847,983
CURRENT FILING DATE: 2004-05-17
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
; OTHER INFORMATION: US-10-425-115-320203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: beta-1,4 N-acetylgalactosaminyl (GalNac)
; OTHER INFORMATION: transferase from C. jejuni strain OH4384 (ORF 5a
; OTHER INFORMATION: of lipooligosaccharide (LOS) biosynthesis locus)
US-10-847-983-17
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                                                              CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
CURRENT FILING DATE: 2003-04-28
RUMBER OF SEQ ID NOS: 369326
SEQ ID NO 320203
LENGTH: 385
TYPE: PRT
ORGANISM: Zea mays
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SEQ ID NO 17
LENGTH: 347
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                                                                                                                                                                                                                             APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Gilbert, Michel
                                             FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
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                    Clone ID: MRT4577_55091C.1.pep
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71.4%;
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Pred. No. 7.8e
2; Mismatches
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Pred. No. 7.8e+02;
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. 7.8e+02;
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Best Local Similarity
Matches 6; Conserva
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APPLICANT: Liu, Jingdong
APPLICANT: Zhou, Yihua
APPLICANT: Kovalic, David K.
APPLICANT: Kovalic, David K.
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Tabaska, Jack E
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53313)B
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 73128
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Publication No. US20040216190A1

GENERAL INFORMATION:
APPLICANT: KOVALIC, David K.

TITLE OF INVENTION: NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH
TITLE OF INVENTION: PLANTS AND USES THEREOF FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53377)B
CURRENT APPLICATION NUMBER: US/10/739,930

CURRENT FILING DATE: 2003-12-18

NUMBER OF SEQ ID NOS: 11088
                                                                                                                                                                                            SEQ ID NO 49028
LENGTH: 392
TYPE: PRT
ORGANISM: Glycine max
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LENGTH: 385
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Publication No. US20040034888A1
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                                                                     Matches
                                                                                    Query Match
Best Local
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Local Similarity 85.7%;
les 6; Conservativo
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 195 YSSGLHS 201
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                                                                                      Similarity
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                                                                                    Score 31; DB 15;
Pred. No. 8.8e+02;
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Pred. No. 8.6e+02;
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Pred. No.
                                                                       Mismatches
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TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement; FILE REFERENCE: 38-21(5323)B; CURRENT APPLICATION UNMBER; US/10/424,599; CURRENT FILING DATE: 2003-04-28; NUMBER OF SEQ ID NOS: 285684

SEQ ID NO 285570

LENGTH: 442

TYPE: PRT
                                                                                       RESULT 38
US-10-425-115-192258
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US-10-424-599-285570
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CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID.NOS: 73128
SEQ ID NO 64369
                                Sequence 192258, Application US/10425115
Publication No. US20040214272A1
GENERAL INFORMATION:
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APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
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ORGANISM: Zea mays
FEATURE:
                                                                                                                                                                                                                                                                                                                        PEATURE: OTHER INFORMATION: Clone ID: PAT_MRT3847_9989C.1.pep
                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Glycine max
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: Kovalic, David K.
: Screen, Steven E
: Tabaaka, Jack E
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85.7%;
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Pred. No. 9.9e+02;
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Pred. No. 9.1e+02;
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Sequence 242547, Application US/10425115;
publication No. US20040214272A1

GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua

APPLICANT: Zhou, Yihua

APPLICANT: Cao, Yongwei

TITLE OF INVENTION: Nucleic Acid Molecules and Oth

TITLE OF INVENTION: Plants

FILE REFERENCE: 38-21(53222)B

CURRENT FAILIG DATE: 2003-04-28;
NUMBER OF SEQ ID NOS: 369326

SEQ ID NO 242547
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US-10-424-599-231545
; Sequence 231545, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
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TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21 (5322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 192258
LENGTH: 501
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 231545
LENGTH: 529
TYPE: PRT
ORGANISM: Glycine max
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Best Local Similarity
Matches 6; Conserv
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Best Local Similarity
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APPLICANT: Kovalic David K
APPLICANT: Zhou Yihua
APPLICANT: Cao Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
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ORGANISM: Zea mays
FEATURE:
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OTHER INFORMATION: Clone ID: PAT_MRT3847_51106C.1.pep
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Pred. No. 1.1e+03;
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Query M Best Lo Matches	RESULT 2 G86895 hypothetical pr C;Species: Lact C;Date: 23-Mar- C;Accession: G8 R;Bolotin, A; Genome Res. 11, A;Title: The co A;Reference num A;Accession: G8 A;Status: preli A;Molecule type A;Residues: 1-3 A;Cross-referen	Query M Best Lo Matches Qy Db	RESULT 1 T30122 C;Species: Ca C;Date: 15-Oc C;Accession: R;Langston, R;Langston, S;Description A;Description A;Description A;Accession: A;Accession: A;Status: pre A;Molecule ty A;Residues: 1 A;Cross -refer A;Experimenta C;Genetics: A;Gene: CESP: A;Gene: CESP: A;Map positio A;Introns: 47		444443333333 5443210
y Match Local Similarity hes 6; Conserv	RESULT 2 G86895 hypothetical protein yweB [impothetical protein groups actis C;Decies: Lactococcus lactis C;Date: 23-Mar-2001 #sequence, C;Accession: G86895 R;Bolotin, A.; Wincker, P.; M Genome Res. 11, 731-753, 2001 A;Title: The complete genome a, Reference number: A86625; M A;Accession: G86895 A;Residues: preliminary A;Rolecule type: DNA A;Residues: 1-368 <sto> A;Cross-references: UNIPROT:Q: A;Experimental source: strain C;Genetics: A;Gene: yweB</sto>	Match Local Similarity Local Similarity 1 YSSNLHS 7 1               22 FSSNLHS 2	proteinenorhal t-1999 T30122 T30122 T30122 T30122 The EMM :: The EMM :: The IMM :: The I		29 78. 29 78.
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B 2; Len ; B 0;	us lactis 01 #text_ 0.; Mala tic acid 11337471 piD:g127	DB 2; Length 11; hes 0; Ind	elegans 1999 #text 1996 id F22H10. /DDBJ /DDBJ 5; PIDN:AA	Ω	
368; els 0;	. 1a 09- ium	219; els 0;	10		Ig kappa chain V r ribosomal protein protein TOBA9 2 (i transcription fact transcription fact transcription fact transcription fact hypothetical prote hypothetical prote plucose-6-phosphat unknown protein, 2 hypothetical prote molybdopterin bios hypothetical prote
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	IL1403) .; Ehrlic estis est		CESP: P2;		

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K; LASINDTOOK, C.C.; GONZALEZ-BOSCH, C.; Bennett, A.B.
Plant Cell 6, 1485-1493, 1994
A;Title: Two divergent endo-beta-1,4-glucanase genes exhibit overlapping expression
A;Reference number: Z15614; MUID:95086382; PMID:7994180
A;Accession: T06350
                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPROT:Q42872; EMBL:U13055; NID:g531904; PIDN:AAA69909.1; A;Experimental source: strain Castlemart; tissue-type pericarp C;Genetics: A;Gene: Cel2 C;Function:
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C;Species: Capsicum annuum (pepper)
C;Species: 24-Jul-1998 #sequence_revision 24-Jul-1998 #text_change 09-Jul-2004
Ig kappa chain
C;Species: Mus
                                                                                                                                                                                                                                                                                          A;Description: hydrolysis of 1,4-beta-D-glucosidic linkages A;Pathway: cellulose degradation C;Superfamily: Arabidopsis membrane-anchored cellulase KOR C;Keywords: glycosidase; hydrolase; polysaccharide degradati F;1-23/Domain: signal sequence #status predicted <SIG>F;24-489/Product: cellulase 2 #status predicted <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-489 < LAS>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: mRNA
A; Residues: 1-324 <FER>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Plant Mol. Biol. 29, 735-747, 1995
A;Title: Differential ethylene-inducible expression of cellulase in pepper plants.
A;Reference number: S61445; MUID:96128016; PMID:8541500
A;Accession: S61447
A;Status: nucleic acid sequence not shown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Cross-references: UNIPROT: Q43751; EMBL: X83711
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                                                             RESULT 5
                                            A48677
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ellulase (EC 3.2.1.4) Cel2 precursor - tomato;Alternate names: endo-1,4-beta-glucanase;Species: Lycopersicon esculentum (tomato);Date: 30-Apr-1999 #sequence_revision 30-Apr-1999 #text_change 09-Jul-2004;Date: 30-Apr-1999
                                                                                                                                                                                                                                                                                                           Superfamily: Arabidopsis membrane-anchored cellulase KOR; Keywords: glycosidase; hydrolase; polysaccharide degradation; 1-23/Domain: signal sequence #status predicted <SIG>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Status: preliminary; translated
                                                                                                                                                                                                                              Query Match
Best Local
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  V-J region musculus ()
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    (house
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                   (48)
                                                                                                                                                                                                                              86.5%;
85.7%;
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mouse)
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                     mouse
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Pred. No.
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C;Acccc.
F; Holroyd, S.; Jagels, K.;
Holroyd, S.; Jagels, K.;
Holroyd, S02-506, 2000
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R;Tassignon, J.; Brait, M.; Jamila, I.; Urbain, J.; Gottlieb, P.; Proc. Natl. Acad. Sci. U.S.A. 90, 9508-9512, 1993  
A;Title: Molecular characterization of monoclonal CRI-A-positive A;Reference number: A48677; MUID:94022404; PMID:8415731  
A;Accession: A48677
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A;Cross-references: UNIPROT:Q9K0V4; GB:AE002401; GB:AE002098; NID:g7225669; PIDN:AAF4089; A;Experimental source: serogroup B, strain MC58
C;Genetics:
C;Genetics:
A;Gene: NMB0455
C;Superfamily: Bacillus conserved hypothetical protein ypsC
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C;Superfamily: immunoglobulin V region; immuno
C;Keywords: immunoglobulin
F;16-90/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                       C;Species: Neisseria meningitidis
C;Date: 05-May-2000 #sequence revision 05-May-2000 #text_change 09-Jul-2004
C;Accession: A81833
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A81833
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A; Molecule type: DNA
A; Residues: 1-380 < PAR>
                         A;Status: preliminary A;Molecule type: DNA
                                                                 A;Title: Complete DNA sequence A;Reference number: A81775; MUI A;Accession: A81833
                                                                                                                                                                                                                                                                  conserved hypothetical protein NMA2030 [imported] - Neisseria meningitidis
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Best Local S
Matches 5
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Best Local
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                                                                                                                                                       Achtman, M.; James, K.D.; Bentley, Jagels, K.; Leather, S.; Moule, S.;
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                                                                                          nce of a serogroup A strain of MUID:20222556; PMID:10761919
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Pred. No.
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Pred. No.
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13;
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Rajandream,
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A;Experimental source: serogroup A, C;Genetics:

A;Cross-references: UNIPROT:Q9RQW4; GB:AL162757; GB:AL157959; NID:g7380371; PIDN:CAB8524: A;Experimental source: serogroup A, strain Z2491

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th, T.; Connerton, P.; Cronin, A.; Davis, P.; Davies, R.M.; Dow S.; Moule, S.; O'Gaora, P.
Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.;
A;Title: Complete genome sequence of a multiple drug resistant A;Reference number: AB0502; MUID:21534947; PMID:11677608
A;Accession: AF0777
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Species: Nostoc sp. PCC 7120
A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. strain PCC C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change 09-Jul-2004 C;Accession: AI2054
C;Accession: AI2054
R;Kaneko, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Notanako, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Notanako, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Notanako, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Notanako, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, Watanabe, S.; Wa
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th. T.; Connerton, P.; Cromin,
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A; Residues: 1-290 < KUR>
A; Cross-references: UNI
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DNA Res. 8, 205-213, 2001
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A; Residues: 1-243 < PAR>
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C;Superfamily: Bacillus conserved hypothetical protein ypsC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: AB1807;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hypothetical protein all1991 [imported] - Nostoc sp. (strain PCC 7120)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Accession: AI2054
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Best Local S
Matches 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cross-references: GB:AL513382;
                                                                                                                                                                                                                                                                                                                                                                         Cross-references: UNIPROT:Q8YVI7; GB:BA000019; PIDN:BAB73690.1; PID:g17131081; GSPDB:(
;Experimental source: strain PCC 7120
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5; Conserv
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                                       YSNNLH 195
                                                                                                      YSSNLH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                S.; Sugimoto, M.; Takazawa, M.; Yamada, M.; Yasuda, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           G.; James,
Cronin, A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               83.8%;
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                                                                                                                                                                                                          81.1%;
83.3%;
                                                                                                                                                                      Score 30; DB Pred. No. 65; Nismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 30; DB 2; Length 243; pred. No. 53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PIDN:CAD02540.1; PID:g16503401; GSPDB:GN00176
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Pred. No.
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Query Match
Best Local Similarity
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C;Genetics: <SCH1>
A;Gene: B24M22.250
A;Map position: 6
A;Introns: 67/3; 320/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: DNA
A;Residues: 1-313 <BEV>
A;Cross-references: UNIPROT:Q9LZ82; EMBL:AL162875
A;Experimental source: cultivar Columbia; BAC clone T32M21
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probable RNA-binding protein - Arabidopsis thaliana
N;Alternate names: protein T32M21.30
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 09-Jul-2004
C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 09-Jul-2004
C;Accession: T44839
R;Bevan, M.; Terryn, N.; Ardiles, W.; Buysshaert, C.; Dasseville, R.; De Clewes, H.W.; Rudd, S.; Lemcke, K.; Mayer, K.F.X.
A;Description: may play a role as A;Note: induced by blue light C;Keywords: mitochondrion
                                                                             A; Introns:
C; Function:
                                                                                                                                                                                                                                                    A;Genetics: SCH1
C;Genetics: <BRU1>
                                                                                                                                                                                                                                                                                                  A;Residues: 1-412 <SCH>
A;Residues: 1-412 <SCH>
A;Cross-references: EMBL:AL390354; PIDN:CAB99393.1; GSPDB:GN00116; NCSP:B24M22.250
A;Experimental source: strain OR74A
                                                                                                                                                                                                                                                                                                                                                                          A;Status: preliminary A;Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                           A; Reference number: Z25286
A; Accession: T51229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Schulte, U.; Aign, V.; submitted to the Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mol. Gen. Genet. 252, 223-229, 1996
A;Title: bli-4, a gene that is rapidly induced by blue light, encodes a novel mitochondr:
A;Reference number: S72617; MUID:96439828; PMID:8842141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Bruchez, J.J.P.; Eberle, J.; Kohler, W.; Kruft, V.; Radford, Mol. Gen. Genet. 252, 223-229, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Date: 24-Jul-1998 #sequence_revision 24-Jul-1998 #text_change 09-Jul-2004 C;Accession: S72617; T51229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           alcohol dehydrogenase homolog Bli-4 [imported] - Neurospora crassa
N;Alternate names: protein B24M22.250
C;Species: Neurospora crassa
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A; Introns: 29/1; 108/3;
A; Note: T32M21.30
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A;Reference number: Z24487
A;Accession: T48439
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A;Experimental_source: mycellum, strain Sta
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       YSONVHS 203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aign, V.; Hoheisel, J.; Brandt, P.; Fartmann, B.; Holland, he Protein Sequence Database, July 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                   (NAD+) -dependent alcohol dehydrogenase
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Score Pred.

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Length 412;

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Nature 408, 816-820, 2000
A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, E.; Kim, A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Maiti, R.; Marzia C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Marzia Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Tallc Ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A;Reference number: A86141; MUID:21016719; PMID:11130712
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A; Introns: 62/2; 97/3;
C; Superfamily: Caenorha
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T23954
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A; Residues: 1-427 <WIL>
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A;Status: preliminary; translated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               submitted to the EMBL Data Library, A; Reference number: Z19823
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
C;Accession: C86247
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A; Residues: 1-424 < STO>
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A;Accession: C86247
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Caenorhabditis elegans hypothetical protein R06B9
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Pred. No. 98;
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war, K.
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                                 A;Title: Genome Sequence of the Plant Pathogen and Biotechnology Agent Agrobacterium
                                                        R;Goodner, B.; Hinkle, G.; Gattung, S.;
A.; Liu, F.; Wollam, C.; Allinger, M.;
Science 294, 2323-2328, 2001
                                                                                                                        C;Species: Agrobacterium tumefaciens C;Date: 30-Sep-2001 #sequence_revision C;Accession: E97624
                                                                                                                                                                                         osmolarity sensor protein envZ (envZ) rp426 [imported] - Agrobacterium tumefaciens
               A; Reference number:
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A97359; MUID:21608551;

PMID:11743194

30-Sep-2001 #text\_change 09-Jul-2004 Miller, N.; Blanchard, M.; Qurol Doughty, D.; Scott, C.; Lappas,

M.; Qurollo, ; Lappas, C.;

B.; Goldn ; Markelz,

Goldman, kelz, B.;

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R.Nolling, J.; Berton, G.; Omelchenko, M.V.; Markarova, K.S.; Zeng, Q.; Gibson, R.; Lee, paly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R.

J. Bacteriol. 183, 4823-4838, 20011
A,Title: Genome Sequence and Comparative Analysis of the Solvent-Producing Bacterium Clos A,Reference number: A96900; MUID:21359325; PMID:21359325
A,Accession: P97296
A,Status: preliminary
A,Accession: Perferences: UNIPROT:Q97E89; GB:AE001437; PIDN:AAK81161.1; PID:g15026298; GSPDB:GNAES: L458 <KUR-A,Cross-references: UNIPROT:Q97E89; GB:AE001437; PIDN:AAK81161.1; PID:g15026298; GSPDB:GNAEXperimental source: Clostridium acetobutylicum ATCC824
C;Gene: CACG2225
C;Superfamily: UDP-N-acetylmuramate-alanine ligase
                                                                                                                                                                                                                                                                                                                                                                                                                             R;Wood, D.W.; Setubal, J.C.; Kaul, R.; Monks, D.; Chen, L.; erage, G.; Gillet, W.; Grant, C.; Guenthner, D.; Kutyavin, T.; Karp, P.; Romero, P.; Zhang, S.
Science 294, 2317-2323, 2001
A;Authors: Yoo, H.; Tao, Y.; Biddle, P.; Jung, M.; Krespan,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   UDP-N-acetylmuramate-alanine ligase [imported] - Clostridium acetobutylicum C;Species: Clostridium acetobutylicum C;Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 09-Jul-2004 C;Accession: F97296
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A;Accession: AF2847
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          two component sensor kinase envz [imported] - Agrobacterium tumefaciens (strain C;Species: Agrobacterium tumefaciens C;Date: 11-Jan-2002 #sequence_revision 11-Jan-2002 #text_change 09-Jul-2004 C;Accession: AF2847
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AF2847
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A;Experimental source: strain C58 (
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100.0%; Pred. No.
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1; Mismatches
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C;Species: Homo sapiens (man)
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999
C;Accession: T17257
R;Koehrer, K.; Beyer, A.; Mewes, H.W.; Gassenhuber, submitted to the Protein Sequence Database, Septemb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Kaneko, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, A.; Iriguchi Nakazaki, N.; Shimpo, S.; Sugimoto, M.; Takazawa, M.; Yamada, M.; Yasuda, M.; Tabata, S DNA Res. 8, 205-213, 2001
A;Title: Complete Genomic Sequence of the Filamentous Nitrogen-fixing Cyanobacterium Ana A;Reference number: AB1807; MUID:21595285; PMID:11759840
A;Accession: AD2532
                                                                                                                                                                                                                                              A;Status: preliminary
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-816 < KOE>
A;Cross-references: UNIPROT:Q9UFT2; EMBL:AL117472
A;Experimental source: adult uterus; clone DKFZp586P1422
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A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change
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A; Accession: T17257
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A;Residues: 1-472 <KUR>
A;Crose-references: UNIPROT:Q8ZSG7; GB:AP003602; PIDN:BAB77186.1; PID:g17134628; GSPDB:Q
A;Experimental source: strain PCC 7120
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A; Map position: circular
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A;Residues: 1-459 <KUR>
A;Cross-references: UNIPROT:Q8UDCO; GB:AE007869; PIDN:AAK87950.1; PID:g15157354; GSPDB:Q
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              YSSNFHA 498
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1.1e+02;
0;
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2e+02;
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ches 0;
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                                                                                                                         R;Fowler, A.V.; Smith, P.J.

J. Biol. Chem. 258, 10204-10207, 1983

J. Biol. Chem. 258, 10204-10207, 1983

A;Title: The active site regions of lacZ and ebg beta-galactosidases are homologous.

A;Reference number: A92390; MUID:83290932; PMID:6411710

A;Contents: annotation; active site regions

R;Hall, B.G.; Betts, P.W.; Wootton, J.C.

Genetics 123, 635-648, 1989

A;Title: DNA sequence analysis of artificially evolved ebg enzyme and ebg repressor.

A;Reference number: S09205; MUID:90128218; PMID:2515108
A;Molecule type: DNA
A;Residues: 13-476,'R',478-651,'S',653-660,'P',662-779,'M',780-1037,'T',1039-1042 <HAL>
A;Cross-references: EMBL:X52031; NID:g41307; PIDN:CAA36274.1; PID:g41309
C;Comment: The wild-type enzyme is an ineffective lactase. Two classes of point mutation
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A;Title: Complete genome sequence of F
A;Reference number: A82950; MUID:20437
A;Recession: B83136
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-844 <STO>
A;Cross-references: UNIPROT:Q9HWU4; GB
A;Experimental source: strain PAO1
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                          C;Accession: B83136
R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, Lory, S.; Olson, M.V.
Nature 406, 959-964, 2000
                                                                                                                                                     A;Gene: PA4084
C;Superfamily: outer membrane usher protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    probable fimbrial biogenesis usher protein PA4084 [imported] - Pseudomonas aeruginosa
C;Species: Pseudomonas aeruginosa
C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 09-Jul-2004
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83.3%;
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50; MUID:20437337; PMID:10984043

    Mismatches

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Pred. No.
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Larbig,
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K.; Lim,
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N;Alternate names: beta-D-galactoside galactohydrolase; lactase; phospho-beta-D-ga (;Species: Bscherichia coli (;Species: Bscherichia coli (;Date: 30-Sep-1987 #sequence revision 05-Dec-1997 #text\_change 09-Jul-2004 (;Accession: A65096; A25751; S09206 R;Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.T.; Burland, V.; Riley, A.; Rose, D.J.; Mau, B.; Shao, Y. Science 277, 1453-1462, 1997 Science 277, 1453-1462, 1997 A;Title: The complete genome sequence of Escherichia coli K-12. A;Reference number: A64720; MUID:97426617; PMID:9278503 A;Accession: A65096 A;Cross-references: UNIPROT:P06864; UNIDROT:Q47170; GB:AE0000389; GB:U00096; NID:g1789451; A;Experimental source: strain K-12, substrain MG1655 R;Stokes, H.W.; Betts, P.W.; Hall, B.G. Mol. Biol. Evol. 2, 469-477, 1985 Mol. Biol. Evol. 2, 469-477, 1985 A;Title: Sequence of the ebgA gene of Escherichia coli: comparison with the lacZ gene. A;Reference number: A93056; MUID:88216133; PMID:3939707 A;Accession: A25751 beta-galactosidase (EC 3.2.1.23) alpha chain - Escherichia coli (strain K-12) N;Alternate names: beta-D-galactoside galactohydrolase; lactase; phospho-beta-D-galactosi A;Status: nucleic acid sequence not shown; translation not shown 4;Residues: A; Molecule type: DNA 1-1042 <BLAT>

A; Reference number: A; Accession: S09206

A;Molecule type: DNA A;Residues: 80-476,'R',478-651,'S',653-660,'P',662-674,'CRSWTPAKRS',685-724,'KCASV',730-7 A;Cross-references: GB:X03228; GB:M13700; GB:M13796; NID:g41311; PIDN:CAA26978.1; PID:g41

PID:g41

genes

YSSNLHS

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A;Molecule type: DNA
A;Residues: 1-1042 <HAY>
A;Residues: 1-1042 <HAY>
A;Cross-references: UNIPROT:Q8XAM9; GB:BA000007; PIDN:BAB37381.1; PID:g13363431; GSPDB:d
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C;Accession: E85968
R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner,
                                                                                                                                                                                                                                                          DNA Res. 8, 11-22, 2001
A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7
A;Reference number: A99629; MUID:21156231; PMID:11258796
                                                                                                                                                                                                                                                                                                                         gasawara, N.; Yasunaga,
DNA Res. 8, 11-22, 2001
                                                                                                                                                                                                                                                                                                                                                                                              evolved beta-D-galactosidase alpha subunit [imported] - Escherichia coli (s
C;Species: Escherichia coli
C;Date: 18-Jul-2001 #sequence_revision 18-Jul-2001 #text_change 09-Jul-2004
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A;Cross-references: UNIPROT:Q8XAM9; GB:AE005174; NID:g12517660; PIDN:AAG58209.1; GSPDB:Q
A;Experimental source: strain O157:H7, substrain EDL933
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Escherichia coli
C;Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change
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C;Keywords: glycosidase; hexamer; hydrolase; magnesium
F;413,415,461/Binding site: magnesium (Glu, His, Glu) #status predicted
F;461,502,524/Active site: Glu, Tyr, Glu #status predicted
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A; Map position: 68 min
C; Complex: homohexamer
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                                                                                                                                                                                                                                                                                                                                         T.; Makino, K.; Ohnishi, M.; Kurokawa, K.; Ishii, K.; Yokoyama, K.;
N.; Yasunaga, T.; Kuhara, S.; Shiba, T.; Hattori, M.; Shinagawa, H.
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K.; Apodaca,
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A;Keau, T.D.; Brunham, R.C.; Shen, C.; Gill, S.R.; Heidelberg, J.F.; White, O.; Hickey, C.; Dodson, R.; Gwinn, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McClarty, G.; Salzber Nucleic Acids Res. 28, 1397-1406, 2000

A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39.

A;Reference number: A81500; MUID:20150255; PMID:10684935
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C81601

polymorphic membrane protein B/C family CP0212 [imported] - Chlamydophila pneumoniae polymorphic membrane protein B/C family CP0212 [imported] - Chlamydophila pneumoniae C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae C;Botes: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 11-May-2000 C;Accession: C81601

C;Accession: C81601

R;Read, T.D.; Brunham, R.C.; Shen, C.; Gill, S.R.; Heidelberg, J.F.; White, O.; Hicke, R;Read, T.D.; Brunham, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McClarty, G.; Salzbe, C.; Dodson, R.; Gwinn, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McClarty, G.; Salzbe
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A;Title: Comparison of whole genome sequences of chlamydia pneumoniae J138 A;Reference number: A86491; MUID:20330349; PMID:10871362
A;Accession: H86557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-1723 <ARN>
A;Rrosereferences: UNIPROT:Q9Z812; GB:AE001638; GB:AE001363; NID:g4376819; PIDN:AAD1868(
A;Cross-references: UNIPROT:Q9Z812; GB:AE001638; GB:AE001363; NID:g4376819; PIDN:AAD1868(
A;Experimental source: strain CWL029
C;Genetics:
C;Genetics:
A;Gene: pmp_20
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E72067
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A;Residues: 1-1723 <STO>
A;Residues: 1-1723 <STOO
A;Cross-references: UNIPROT:Q9Z812; GB:BA0000008; NID:g8978911; PIDN:BAA98746.1; GSPDB:GN(
A;Experimental source: strain J138
C;Genetics:
A;Gene: pmp_20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Comparative genomes of Clamydia pneumoniae and C. trachomatis. A;Reference number: A72000; MUID:99206606; PMID:10192388
A;Accession: E72067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polymorphic membrane protein B family - Chlamydophila pneu C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae C;Date: 23-Apr-1999 #text_C;Date: 23-Apr-1999 #text_C C;Accession: E72067 C;Accession: E72067 R;Kalman, S; Mitchell, W.; Marathe, R.; Lammel, C.; Fan, Nature Genet. 21, 385-389, 1999
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Chlamydophila pneumoniae, Chlamydia pneumoniae
Apr-1999 #sequence_revision 23-Apr-1999 #text_change 09-Jul-2004
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J. 4.5e+02;
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4.5e+02;
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SHISSTY

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A;Molecule type: protein
A;Residues: 1-108 <SCH>
A;Cross-references: UNIPROT:P01643
A;Cross-references: UNIPROT:P01643
C;Comment: This chain was isolated from a myeloma protein.
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kaphain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into la C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer
C;Keywords: heterotetramer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ig kappa chain V region (Py69) - mouse (Species: Mus musculus (house mouse) (Species: Mus musculus (house mouse) (Species: Mus musculus (house mouse) (Species: Mus musculus (house) (Species: Mus musculus (Py60) (Species: Gasta) (Species: Gasta)
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A; Residues: 1-111 < RUF>
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A;Status: preliminary; nucleic acid
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Eur. J. Biochem. 59, 525-537, 1975
A;Title: Determination of the prim
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A;Residues: 1-1732 <REA>
A;Cross-references: GB:AE002182; GB:AE002161; NID:g7189140; PIDN:AAF38082.1; PID:g718914
A;Experimental source: strain AR39, HL cells
                                                                                                                               Query Match
Best Local
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Species: Mus musculus (house mouse)
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Ig kappa chain V region (Py20) - mouse (Species: Mus musculus (house mouse) (Species: Mus musculus (house mouse) (C;Date: 18-Oct-1991 #sequence_revision 18-Oct-1991 #text_change 09-Jul-2004 (C;Accession: A38740 R;Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R. J. Biol. Chem. 266, 6607-6613, 1991 A;Biol. Chem. 266, 6607-6613, 1991 A;Title: Heavy and light chain variable region sequences and antibody properties of A;Reference number: A38740; MUID:91177923; PMID:1706720 A;Accession: A38740
                                          R;Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R. J. Biol. Chem. 266, 6607-6613, 1991
A;Title: Heavy and light chain variable region sequences and A;Reference number: A38740; MUID:91177923; PMID:1706720
A;Accession: C38740
                                                                                                                                                              Ig kappa chain V region (Py2) - mouse (;Species: Mus musculus (house mouse) C;Date: 18-Oct-1991 #sequence_revision 18-Oct-1991 #text_change C;Accession: C38740
                                                                                                                                                                                                                                                            RESULT
C38740
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
E;19-93/Domain: immunoglobulin homology <IMM>
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A;Title: Heavy and light chain variable region sequences and A;Reference number: A38740; MUID:91177923; PMID:1706720
A;Accession: E38740
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138740

19 kappa chain V region (Py54) - mouse

C;Species: Mus musculus (house mouse)

C;Date: 18-Oct-1991 #sequence_revision 18-Oct-1991 #text_change

C;Accession: E38740

C;Accession: E38740
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C;Superfamily: immunoglobulin V region; imm
C;Keywords: heterotetramer; immunoglobulin
F;19-93/Domain: immunoglobulin homology <IN
A_iStatus: preliminary; nucleic acid sequence not shown; not compared with conceptual tran A_iMolecule type: mRNA
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A;Molecule type: mRNA
A;Residues: 1-111 <RUF>
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71.4%;
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C;Accessic
R;Burger,
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A; Residues: 1-243 <STO>
                                                                                                                                                                                                                                                                        Science 282, 2012-2018, 1998
A;Title: Genome sequence of the nematode C. elegans: a platform for investigating biolog A;Reference number: A75000; MUID:99069613; PMID:9851916
A;Reference number: A75000; MUID:99069613; PMID:9851916
A;Rote: see websites genome.wustl.edu/gsc/C_elegans/ and www_sanger.ac.uk/Projects/C_ele A;Note: published errata appeared in Science 283, 35, 1999; Science 283, 2103, 1999; and A;Accession: G89567
                                                                                                                                                                                                                                                                                                                                                                                 C;Date: 10-May-2001 #sequence_revision 10-May-2001 C;Accession: G89567 C;Accession: G89567 R;anonymous, The C. elegans Sequencing Consortium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Burger, G.; Saint-Louis, D.; Gray, M.W.; Lang, B.F. submitted to the EMBL Data Library, December 1998 A;Description: Complete sequence of the mitochondrial A;Reference number: Z17255
A;Accession: T11242
A;Status: preliminary; translated from GB/EMBL/DDBJ
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C;Superfamily: immunoglobulin
C;Keywords: heterotetramer; in
F;19-93/Domain: immunoglobulir
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                                Query Match
Best Local Similarity
""-" hes 5; Conserve
                                                                                                                                                                                                                                                        A; Status: preliminary
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A;Residues: 1-227 <BUR>
A;Cross-references: UNIPROT:O99993; EMBL:AF114794; NID:g4106927; PID:g4106954; PIDN:AADC
C;Genetics:
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C;Species: Caenorhabditis elegans
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;Note: similar to C. elegans
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                                                   Score 29; DB Pred. No. 88; 0; Mismatches
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N;Alternate names: transforming protein (jun)
C;Specises: Rattus norvegicus (Norway rat)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change
C;Accession: $12742; $20028; A37381
R;Kitabayashi, I.; Saka, F.; Gachelin, G.; Yokoyama, K.
Nucleic Acids Res. 13, 3400, 1990
A;Title: Nucleotide sequence of rat c-jun protooncogene.
A;Reference number: $12742; MUID:90287724; PMID:2113275
A;Accession: $12742
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C;Superfamily: jun transforming protein; fos/jun DNA-binding domain homology
C;Keywords: DNA binding; leucine zipper; phosphoprotein; proto-oncogene; tran
F;250-290/Domain: fos/jun DNA-binding domain homology <FJD>
F;283-311/Region: leucine zipper motif
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NyAlternate names: fos-associated 39K protein; protein PEA1; transcription:
C;Species: Mus musculus (house mouse)
C;Date: 31-Dec-1989 #sequence revision 31-Dec-1989 #text_change 09-Jul-2004
C;Accession: A31145; S04683; S04537
                                                                                                      A;Cross-references: UNIPROT:P17325; EMBL:X17215; NID:g57079; PIDN:CAA35084.1; | R;Kitabayashi, I.; Kawakami, Z.; Chiu, R.; Ozawa, K.; Matsuoka, T.; Toyoshima, EMBO J. 11, 167-175, 1992
A;Title: Transcriptional regulation of the c-jun gene by retinoic acid and E1A A;Reference number: S20028; MUID:92155155; PMID:1310930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nature 334, 535-537, 1988
A;Title: Transcriptional activation of c-jun during the G(0)/G(1) transition A;Reference number: S04537; MUID:88302446; PMID:3136397
A;Accession: S04537
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Proc. Natl. Acad. Sci. U.S.A. 85, 8464-8467, 1988
A;Tille: Induction of protooncogene c-jun by serum growth factors.
A;Reference number: A31345; MUID:89042204; PMID:3186736
A;Accession: A31345
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A;Residues: 1-182,'C',184-334 <RYS>
A;Cross-references: EMBL:X12761; NID:g52758; PIDN:CAA31252.1;
C;Genetics:
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A; Molecule type:
A; Residues: 1-33:
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A; Residues: 1-334 < KIT>
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A; Residues: 1-334 < RYD>
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                                                   translation not shown
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C;Species: Schizosaccharomyces pombe
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004
C;Accession: T38822
                                                                                                                                                    R;Connor, R.; Churcher, C.M.; Barrell, B.G.; Rajandream, submitted to the EMBL Data Library, September 1997 A;Reference number: Z21813 A;Accession: T38822
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C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
C;Accession: T25192
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C;Superfamily: jun transforming protein; fos/jun DNA-binding domain homology
C;Keywords: DNA binding; leucine zipper; nucleus; phosphoprotein; transcription
F;250-290/Domain: fos/jun DNA-binding domain homology <FUD>
                                        A;Cross-references: UNIPROT:036030; EMBL:Z98980; PIDN:CAB11721.1; GSPDB:GN00066; SPDB:SFA;Experimental source: strain 972h-; cosmid c4F10
                                                                                    A;Status: preliminary; translated from GB/EMBL/DDBJA;Molecule type: DNAA;Residues: 1-391 <CON>
                                                                                                                                                                                                                                                                                                                   hypothetical protein SPAC4F10.18 - fission yeast (Schizosaccharomyces pombe)
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A; Introns: 74/2; 128/3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Reference number: A; Accession: T25192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hypothetical protein T23G11.2 - Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Gene: jun
C;Function:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Title: Structure and expression of the rat c-jun messenger RNA: tissue distribution A;Reference number: A37381; MUID:90002916; PMID:2507134 A;Accession: A37381
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Sakai, M.; Okuda, A.; Hatayama
Cancer Res. 49, 5633-5637, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: EMBL:X17215; NID:g57079; PIDN:CAA35084.1; PID:g57080 R;Sakai, M.; Okuda, A.; Hatayama, I.; Sato, K.; Nishi, S.; Muramatsu, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gene: CESP:T23G11.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Experimental source:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cross-references: GB:X17163; NID:g57819; PIDN:CAA35041.1; PID:g57820
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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SPDB:SPAC4F10.18
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Pred. No. 1.3e
0; Mismatches
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1.3e+02;
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glucose-6-phosphate isomerase pgi - Bacillus subtilis
C;Species: Bacillus subtilis
C;Species: Bacillus subtilis
C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 09-Jul-2004
C;Accession: B69675
C;Accession: B69675
C;Rocession: B69675
R;Kunst, F.; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Berterc, C.; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Chorler, C.; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Chorler, Barrisch, S.D.; Emmerson, P.T.; Entian, K.D.; Exrington, J.; Fabret, C.; Ferrari, E.
Nature 390, 249-256, 1997
A;Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A.; Gallerc, iech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hullo, M.F., Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Lardinois, A.; Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S.; Maueel, Y., Mathors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, S.; Maueel, A.; Winters, P.; Wipat, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; Uchiyama, A.; Authors: Yoshikawa, H.F.; Zumstein, B.; Yoshikawa, H.; Danchin, A.
A;Tile: The complete genome sequence of the Gram-positive bacterium Bacillus subtilis. A; Accession: B69675
A. Steries not shown
unknown protein, 29405-27288 [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Dete: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
C;Accession: 096655
R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Pederspiel, N.A.; Kaul, S.; White, O.; Alonso, Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K.; ansen, N.F.; Hughes, B.; Huizar, L.
Nature 408, 816-820, 2000
A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, B.; Kim, C.; C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Waiti, R.; Marziall, Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sun, H.; Tallon, I ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A;Reference and analysis of Chromosome 1 of the plant Arabidopsis.
A;Accession: 096655
A;Accession: 096655
A;Accession: 096655
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G96655
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C; Superfamily:
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A; Introns: 9/2; 35/1
C; Superfamily: Schiz
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Best Local S
Matches : 5
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Best Local
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5; Conserv
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5; Conserv
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83.3%;
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71.4%;
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Pred. No.
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Pred. No. 1.7e+02
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C;Date: 24-May-2001 #sequence_revision 24-May-2001 #text_change 16-Aug-2004 C;Accession: F90407 C;Accession: F90407 R;She, O; Singh, R.K.; Confalonieri, F.; Zivanovic, Y.; Allard, G.; Awayez, M.J.; Cha Jong, I.; Jeffries, A.C.; Kozera, C.J.; Medina, N.; Peng, X.; Thi-Ngoc, H.P.; Redder, arrett, R.A.; Ragan, M.A.; Sensen, C.W.; Van der Oost, J. submitted to GenBank, April 2001 submitted to GenBank, April 2001 A;Description: Sulfolobus solfataricus complete genome.
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T33061
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A;Introns: 69/2; 144/3; 230/2; 446/3
C;Superfamily: Caenorhabditis elegans hypothetical protein F47H4.4
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                                                                                                                                                                                                        \bar{\mathbf{A}}_i; Cross-references: UNIPROT: Q97W71; GB:AE006641; NID: g13815669; PIDN:AAK42517.1; GSPDB:
                                                                                                                                                                                                                                  A; Molecule type: DNA
A; Residues: 1-550 < KUR>
                                                                                                                                                                                                                                                                                                   A; Reference number: A99139
A; Accession: F90407
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F90407
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-516 <STO>
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A; Accession: T33061
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A; Description: The sequence of C. elegans cosmid F56C3
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C;Species: Caenorhabditis elegans
C;Species: Caenorhabditis elegans
C;Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change 09-Jul-2004
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A; Residues: 1-465 < STO>
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Best Local S
Matches 5
                                                                                                                                   :Superfamily: Molybdenum cofactor molybdenum incorporation protein MoeA with molybdate
                                                                                                                                                                Gene: moeA-2
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Best Local Similarity 83.3
Matches 5; Conservative
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nes 5; Conserv
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rative 0; Mismatches
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Db 208 YESNLH 213

Search completed: December 17, 2004, 18:30:16 Job time: 3.75281 secs

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Result
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Maximum Match 100%
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OM protein - protein search,
    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
      Query
Match
   BLOSUM62
Gapop 10.0 , Gapext 0.5
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192.513 Million cell updates/sec
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1: uniprot_sprot:*
2: uniprot_trembl:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
    GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BB
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 Q9FCO3
Q62788
Q94192
Q9CDQ1
Q8RRQ4
Q8A9TO
Q9A9TO
Q9F1Y9
Q9F1Y9
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Q9F1Y9
Q9FAV1
Q9FAV1
Q9FAW1
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Q43751
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Q7RIQ6
SKIP MOUSE
AAH66112
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<b>4</b> 5	44	43	42	41	40	39	38	37	36	35	34	မ	32
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83.8	83.8	83.8	83.8	83.8	83.8	83.8	83.8	86.5	86.5	86.5	86.5	86.5	86.5
380	380	347	347	347	347	347	335	615	598	536	500	489	485
N	N	N	N	N	N	N	N	N	N	N	N	N	N
Q9K0V4	Q9RMM0	AAR82873	Q938X8	Q9F0N1	Q9LAK4	Q9L9Q6	Q8RFN2	Q9RP84	Q7Q8M6	Q83Z58	Q93WZ0	Q42872	Q96547
Q9k0v4 neisseria m	Q9rmm0 neisseria m	Aar82873 campyloba	Q938x8 campylobact	Q9f0n1 campylobact	Q9lak4 campylobact	Q919q6 campylobact	Q8rfn2 fusobacteri	Q9rp84 acholeplasm	Q7q8m6 anopheles g	Q83z58 mesoplasma	Q93wz0 nicotiana t	Q42872 lycopersico	Q96547 capsicum an

## ALIGNMENTS

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SEQUENCE FROM N.A.  STRAIN-17XNI;  PubMed=12368865;  Carlton J.M., Angiuoli S.V., Suh B.B., Kooij T.W., Pertea M.,  Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Koo H.L.,  Silva J.C., Ermolaeva M.D., Shumway M.F., Bidwell S.L.,  Peterson J.D., Pop M., Kosack D.S., Shumway M.F., Bidwell S.L.,  Shallom S.J., van Aken S.E., Riedmuller S.B., Feldblyum T.V.,  Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M.,	IT 2  ORTIQ6 PRELIMINARY; PRT; 404 AA.  OTRIQ6; O1-MAR-2004 (TrEMBLrel. 26, Created) O1-MAR-2004 (TrEMBLrel. 26, Last sequence update) O1-MAR-2004 (TrEMBLrel. 26, Last annotation update) Asparagine-rich protein. Name-PY03561; Plasmodium yoelii yoelii. Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium. NCBI_TaxID=73239; [1]	k/DDBJ databases.  NKNOWN_1.  1F83657D44 CRC64;  DB 2; Length 4780 2.8e+02; hes 0; Indels	BIE54 PRELIMINARY; PRT; 4780 A BIE54;  1-MAR-2003 (TYEMBLrel. 23, Created)  1-MAR-2003 (TYEMBLrel. 23, Last sequence  1-MAR-2003 (TYEMBLrel. 26, Last annotatic  ypothetical protein PP13_0148.  ame=PF13_0148;  blasmodium falciparum (isolate 3D7).  karyota; Alveolata; Apicomplexa; Haemosp  CBI_TaxID=36329;  arris B., Lennard N., Clark L., Line A.,  arris B., Lennard N., Clark L., Line A.,

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           REPLINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;

WEDLINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;

WA MIRITINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;

WA Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kondo S.,

WA Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiyosawa H.,

WA Nikaido I., Osato N., Saito R., Schonbach C., Gojobori T.,

WA Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,

WA Schriml L.M., Kanapin A., Matsuda H., Batalov S., Beisel K.W.,

WA Schriml L.M., Eradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,

WA Schriml L.M., Eradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,

WA Schriml L.M., Eradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,

WA Schriml G., Braticolh N., Gissi C., Godzik A., Gough J.,

WA Gasterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,

WA Gasterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,

WA Gasterland T., Gariboldi M., Gissi C., Godzik A., Gough J.,

WA Gasterland T., Kawasawa Y., Kedzierski R.M., King B.L.,

WA Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,

WA Kanai A., Kawaji H., Kawasawa Y., Kedzierski R.M., King B.L.,

WA Magalott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,

WA Nagashima T., Numata K., Okido T., Pavan W. J., Pertea G., Pesole G.,

Petrovsky N., Pillai R., Pontius J.U., Oi D., Ramachandran S.,

Petrovsky N., Pillai R., Pontius J.U., Ring B.Z., Ringwald M.,

Parkan Magalott D., Reid J.C., Reed D.J., Reid J., Ring B.Z., Ringwald M.,

Parkan Magalott D., Reid J.C., Reed D.J., Reid J., Ring B.Z., Ringwald M.,

Parkan Magalott D., Reid J.C., Ringwald M.,

Parkan Magalott D., Reid J.C., Ringwald M.,
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Best Local S
Matches 6
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QBC516; O09040;
10-OCT-2003 (Rel. 42, Created)
29-MAR-2004 (Rel. 43, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Genome sequence and comparative analysis of the model rodent malaria parasite Plasmodium yoelii yoelii.";
Nature 419:512-519(2002).
-i- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cunningham D.A., Preiser P.R van Lin L.H., Janse C.J., Wa van Lister S.L., Venter J.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hamilton B.A., Smith D.J., Mueller K.L., Kerrebrock A.W., Bronson R.T., van Berkel V., Daly M.J., Kruglyak L., Reeve M.P., Nemhauser J.L., Hawkins T.L., Rubin E.M., Lander E.S.; "The vibrator mutation causes neurodegeneration via reduced expression of PITP alpha: positional complementation cloning and extragenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOUSE
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Beeson M., Gordon L., Be
Submitted (FEB-2002) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN=ILS, and ISS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
MEDLINE=97325785; PubMed=9182797;
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Mus musculus (Mouse).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neuron 18:711-722(1997).
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; AABL01001031; E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6
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Rodentia;
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47311 MW; '
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bennett
                                                                                                                                                                                                                                                                                                                                                                                                                                     J., Conroy O., Xu Y., Yang F., Canniff J., ennett B., Johnson T.E., Sikela J.M.; the EMBL/GenBank/DDBJ databases.
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Waters A.P., Smith H.O., White O.R.,
.., Fraser C.M., Hoffman S.L., Gardner M.J.,
Semple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 34; DB Pred. No. 77; 1; Mismatches
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RESULT 4
AAH66112
ID AAH6
AC AAH6
DT 14-A

AAH66112 AAH66112;

14-APR-2004

PRELIMINARY; (TrEMBLrel.

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PRT;

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Best Local :
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EMBL; U96726; AAC60757.1; --
EMBL; AF483522; AAL90797.1; --
EMBL; AF483523; AAL90797.1; --
EMBL; AK05436; BAC35778.1; --
EMBL; AK078104; BAC37726.1; --
EMBL; AK078104; BAC37126.1; --
HSSP; O43001; 1192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sultana R., Takenaka Y., Taylor M.S., Teasdale R.D., Tomita M., Verardo R., Wagner L., Wahlestedt C., Wang Y., Watnabe Y., Wells Wilming L.G., Wynshaw-Boris A., Yanagisawa M., Yang I., Yang I., Yang Z., Zavolan M., Zhu Y., Zimmer A., Carninci P., Hayatsu N., Hirozane-Kishikawa T., Konno H., Nakamura M., Sakazume N., Sato Y. Shiraki T., Waki K., Kawai J., Aizawa K., Arakawa T., Fukuda S., Hara A., Hashizume W., Imotani K., Ishii Y., Itoh M., Kagawa I., Miyazaki A., Sakai K., Sasaki D., Shibata K., Shinagawa A., Yasunishi A., Yoshino M., Waterston R., Lander B.S., Rogers J., Hirray R., Harashizari V.
                                                                                                                                                                                                                                                                     888888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Birney E., Hayashizaki Y.;
"Analysis of the mouse transcriptome based on functional annotation 60,770 full-length cDNAs.";
Nature 420:563-573(2002).
                                                                                                                                                                                                      InterPro; IPR005135; Exo
InterPro; IPR000300; IPPC
Pfam; PF03372; Exo_endo_p
SMART; SM00128; IPPC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <del>:</del>
                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -
                                                                                                                                           CONFLICT
                                                                                                                                                                         Endoplasmic reticulum; Hydrolase
DOMAIN 34 337 Cata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Enterpean Bioinformatics Institute. There are no restitute by non-profit institutions as long as its content
                                                                                                                                                          NIAMOC
                                                                                                                                                                                                                                                                   GO:0005829; C:cytosol; ISS.
GO:0043005; C:neuronal cell projection; ISS.
GO:0016312; F:inositol bisphosphate phosphatase activity; ISS.
GO:0046302; F:inositol trisphosphate phosphatase activity; ISS.
GO:0046577; F:lipid phosphatase activity; ISS.
GO:0042577; F:lipid phosphatase activity; ISS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      trisphosphate. May negatively regulate assembly of the actin cytoskeleton (By similarity).

CATALYTIC ACTIVITY: D-myo-inositol 1,4,5-trisphosphate + H(2)0 myo-inositol 1,4-bisphosphate + phosphate.

SUBCELIULAR LOCATION: Endoplasmic reticulum. Following stimulativity with EGF translocates to membarae ruffles (By similarity).

SIMILARITY: Belongs to the inositol-1,4,5-trisphosphate 5-phosphatase type II family.
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434
                             1 YSSNLHS
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                                                                            Similarity
                                                                                                                         468 AA;
                                                             Conservative
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AAC60757.1; -.
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                                                                            91.9%;
85.7%;
                                                                                                                                                                                                                                       IPPC
                                                                                                                                                                                                                                                      endo phos.
                                                                                                                           MW:
                                                                            Score 34;
Pred. No.
                                                                                                                           Required for ruffle localization D -> E (in Ref. 3; BAC37126).; F2E1CA370B97A8A1 CRC64;
                                                                                                                                                                           Catalytic
                                                                                                                                                                                                                          ۲.
                                                               Mismatches
                                                                                                                                                                                                                                                                       organization and biogenesis; ISS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               moved. Usage by and for commercial (See http://www.isb-sib.ch/announce/
                                                                          рв
91;
                                                                                                                                                                         (Potential)
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                                                                                             Length 468
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abe Y., Wells C.,
                                                                                                                                                                                                                                                                                                        activity; ISS
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su N.,
Sato K.,
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RX Strausberg R.L., Feingold B.A., Grouse L.H., Derge J.G.,

RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschanko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

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RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Fonshiyuki S., Carninci P., Frange C.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Frange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Mozny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

VA Hitling M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

VA Holton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whitling M., Maddan A., Young A.C., Shevchenko Y., Bouffard G.G.,

VA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

Nordes G. J. Mayra M. A.,

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Best Local S
Matches 6
SEQUENCE FROM N.A.
STRAIN-A3(2) / M145;
STRAIN-A3(2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OSPCO3;
01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update
01-JUN-2003 (TrEMBLrel. 24, Last annotation upda
Putative secreted amidase.
OrderedLocusNames=SCO7179; ORFNames=SCBA11.07c;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; BCC
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteria, Actinobacteria, Actinobacteridae, Actinomycetales, Streptomycineae, Streptomycetaceae, Streptomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
STRAIN=C57BL/6; TISSUE=Brain;
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=1902;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Streptomyces coelicolor.
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14-APR-2004
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ubmitted (FEB-2004) to the EMBL/GenBank/DDBJ databases
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VCE 468 AA; 54158 MW;
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7; PubMed=12477932;
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27,
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Pred. No. 91;
1; Mismatches
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                                                                                                                                                                   InterPro; IPR001909; KRAB:
InterPro; IPR007087; Znf C2H2.
Pfam; PP001352; KRAB; 1.
Pfam; PF00096; zf-C2H2; 19.
ProDom; PD000003; Znf C2H2; 17.
SMART; SM00349; KRAB; 1.
SMAXT; SM00349; KRAB; 1.
SMO31TE; PS50805; KRAB; 1.
PROSITE; PS00028; ZINC FINGER C2H2_1; 17.
PROSITE; PS00157; ZINC FINGER C2H2_1; 11.
PROSITE; PS00157; ZINC FINGER C2H2_1; 11.
PROSITE; PS00157; ZINC FINGER C2H2_2; 11.
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Hopwood D.A.;
"Complete genome sequence of
coelicolor A3(2).";
Nature 417:141-147(2002).
EMBL; AL393130; CAC01579.1;
HSSP; P82974; 1J3G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  [1]
SEQUENCE FROM N.A.
MEDLINE=96033674; PubMed=7595478;
Pott U., Thiesen H.J., Colello R.J., Schwab M.E.;
Pott U., Thiese zinc finger gene, rKr2, is expression of the color 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q62788
Q62788;
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01-NOV-1996 (TrEMBLrel. 01, La
01-MAR-2004 (TrEMBLrel. 26, La
Cy82/Hi82 zinc finger protein.
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HSSP; P25490;
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                                                                                                                                                                                                                                                                                                                                                                                                                      GO; GO:0005634; C:nucleus; IEA.
GO; GO:0003676; F:nucleic acid binding;
GO; GO:0008270; F:zinc ion binding; IEA.
GO; GO:0006355; P:regulation of transcri
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Mammalia; Eutheria; Rodentia;
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                                                                                       Conservative
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Q94192;
Q9-FEB-1997
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                  MEDIINE-21235186; PubMed=11337471;
Bolotin A., Wincker P., Mauger S., Jaillon O., Malarme K.,
Weissenbach J., Ehrlich S.D., Sorokin A.;
"The complete genome sequence of the lactic acid bacterium Lactococcus
lactis ssp. lactis IL1403.";
Genome Res. 11:731-753(2001).
EMBL, AE006445; AAK06265.1; -.
                                                                                                                                                                                                                                                                         Lactococcus lactis (subsp. lactis) (Streptococcus lactis). Bacteria; Firmicutes; Lactobacillales; Streptococcaceae; LNCBI_TaxID=1360;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (SEP-2003) to the EMBL/GenBank/DDBJ EMBL; U7084; AAB09103.1; -. PIR; T30122; T30122.

WormPep; F32H10.6; CE09575.

Hypothetical protein.
SEQUENCE 219 AA; 25258 MW; 2091CC121B3E5FI
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STRAIN=Bristol N2;
MEDLINE=99069613; PubMed=9851916;
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                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                      Hypothetical protein yweB.
Name=yweB; OrderedLocusNames=LL2167;
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Submitted (OCT-1996)
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Pred. No. 63;
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RA Matsuo Y., Suzuki M., Kasai H., Shizuri Y., Harayama S.;
RA Matsuo Y., Suzuki M., Kasai H., Shizuri Y., Harayama S.;
RA Matsuo Y., Suzuki M., Kasai H., Shizuri Y., Harayama S.;
RA Matsuo Y., Suzuki M., Kasai H., Shizuri Y., Harayama S.;
RI Silver M., Shizuri Y., Harayama S.;
RT "Isolation and phylogenetic characterization of bacteria capable of inducing differentiation in the green alga Monostroma oxyspermum.";
RT FUNCTION: DNA gyrase negatively supercoils closed circular double-cranded DNA in an ATP-dependent manner and also catalyzes the interconversion of other topological isomers of double-stranded DNA interconversion of other topological isomerse family.

CC -I- SIMILARITY: Belongs to the type II topoisomerase family.

EMBLI AB073038; BAB88340.1; -.

DR GO; GO:0003524; F:ATP binding; IEA.

GO; GO:0003524; F:NA binding; IEA.

GO; GO:0003525; F:NA topologomerase (ATP-hydrolyzing) activity; IEA.

GO; GO:000525; P:DNA topologomerase (ATP-hydrolyzing) activity; IEA.

GO; GO:000525; P:DNA topologomerase (ATP-hydrolyzing) activity; IEA.
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SMART; SM00433; TOP2c; 1
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Pfam; PF02518; HATPase_c; 1.
PRINTS; PR00418; TPI2FAMILY.
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InterPro; IPR001241; DNA_topoisoII.
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391 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          T-588.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         43486 MW;
                                                                                                                                                                                                                                                                                                                                                          89.2%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          89.2%;
85.7%;
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21, Last sequence update)
25, Last annotation updat
C 5.99.1.3) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Topoisomerase.
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                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                             Score 33;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 33;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l protein.
C1243C3B9F140D0D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                         OC5E029383468759 CRC64;
                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
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                                                                                                                                                                                                                                                                                                                                                             .2e+02
                                                                                                                                                                                                                                                                                                                                                                                   Length 391;
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                                                                                                                                                                                                                                                                                                                                   <u>,</u>
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Q845X4; 01-JUN-2003 01-JUN-2003

(TrEMBLrel. PRELIMINARY;

24,

Created) Last seq

sequence update)

PRT;

470

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Q845X4

01-MAR-2004 (TrEMBLrel.

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RESULT 11
Q9AJIO
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C: -: SIMILARITY: Belongs to the type II topoisomerase family.

REMBL; AB071140; BAC65173.1; -.

RGO; GO:0005524; F:ATP binding; IEA.

RGO; GO:0003677; F:DNA binding; IEA.

RGO; GO:0003677; F:DNA binding; IEA.

RGO; GO:0003918; F:DNA topoisomerase (ATP-hydrolyzing) activity;

RGO; GO:0006265; P:DNA modification; IEA.

RGO; GO:0006265; P:DNA modification; IEA.

RGO; GO:0006265; P:DNA modification; IEA.

RINTERPO; IPRO01541; ATPbind_ATPase.

RINTERPO; IPRO01541; DNA topoisoII.

RINTERPO; IPRO01541; DNA topoisoII.

RINTERPO; IPRO016171; Toprim_dom.

RFIAM; PF001751; Toprim_dom.

RFIAM; PF001751; Toprim; 1.

RRINTS; PR00418; TP12FAMILY.

RRINTS; PR00418; TP12FAMILY.

RRINTS; PR00418; TP12FAMILY.
Suzuki M., Takadera T., Harayama S., Yamamoto S.;
Submitted (AUG-2000) to the EMBL/GenBank/DbBJ databases.
-i- FUNCTION: DNA gyrase negatively supercoils closed circular double-
stranded DNA in an ATP-dependent manner and also catalyzes the
interconversion of other topological isomers of double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q9AJIO;
                                                                                                                                                                                                                                                                                                                                                                        01-JUN-2001 (TrEMBLrel. 17, 01-JUN-2001 (TrEMBLrel. 17, 01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NON_TER
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Matsuo Y., Suzuki M., Kasai H., Shizuri Y., Harayama S.;
"Isolation and phylogenetic characterization of bacteria capable inducing differentiation in the green alga Monostroma oxyspermum. Environ. Microbiol. 5:25-35(2003).
-I- FUNCTION: DNA gyrase negatively supercoils closed circular do
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ProDom; PD149633; DNA gyrase_B; 1.
SMART; SM00433; TOP2c; 1.
                                                                                                                                                   STRAIN-MBIC4478;
                                                                                                                                                                                                                                                                marine CFB-group bacterium MBIC04478 Bacteria; Bacteroidetes.
                                                                                                                                                                                                                                                                                                                       Name=gyrB;
                                                                                                                                                                                                                                                                                                                                                 DNA gyrase B subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQÜENCE
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NCBI_TaxID=76595;
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                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                   NCBI_TaxID=135558;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FUNCTION: DNA gyrase negatively supercoils closed circular double-
stranded DNA in an ATP-dependent manner and also catalyzes the
interconversion of other topological isomers of double-stranded
DNA rings, including catenanes and knotted rings (By similarity).
CATALYTIC ACTIVITY: ATP-dependent breakage, passage and rejoining
of double-stranded DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   YSENLHS 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       YSSNLHS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PS00177; TOPOISOMERASE_II; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteroidetes; Flavobacteria;
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470 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRELIMINARY;
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85.7%;
                                                                                                                                                                                                                                                                                                                                              (Fragment)
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                                                                                                                                                                                                                                                                                                                                                                     Last sequence update)
Last annotation update)
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Pred. No. 1.5e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        474
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PRESIDENT PRESID
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Matches 6
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cc of SIMILARITY: Belongs to the type II topoisomerase family.

R GO; GO:0005524; F:ATP binding; IEA.

gO; GO:0003677; F:DNA binding; IEA.

gO; GO:0003918; F:DNA topoisomerase (ATP-hydrolyzing) activity;

R GO; GO:0006304; P:DNA modification; IEA.

gO; GO:0006305; P:DNA modification; IEA.

R InterPro; IPR001594; ATPbind_ATPase.

InterPro; IPR001594; ATPbind_ATPase.

InterPro; IPR001541; DNA topoisoII.

InterPro; IPR001741; DNA topoisoII.

Pfam; PP02518; HATPase C; 1.

R Pfam; PP02518; HATPase C; 1.

R Pfam; PP03751; Toprim; 1.

R Pfam; PD03751; Toprim; 1.
DNA rings, including catenanes and knotted rings (By sim catenanes and knotted rings (By sim catenanes and knotted rings) (By sim catenanes and knotted rings) (By sim catenanes and catenanes (By GO:0005524; F:ATP binding; IEA.

R GO; GO:0003677; F:DNA binding; IEA.

R GO; GO:0003918; F:DNA topoisomerase (ATP-hydrolyzing) activing GO: GO:0006304; P:DNA modification; IEA.

R GO; GO:0006265; P:DNA topoisomerase; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q9ETS8 PRELIMINARY;
Q9ETS8;
Q1-MAR-2001 (TrEMBLrel. 16,
01-MAR-2001 (TrEMBLrel. 16,
05-JUL-2004 (TrEMBLrel. 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=IF015992, and IAM14318;
MEDLINE=21477783; PubMed=11594591;
MIDIANE=21477783; PubMed=11594591;
MIDIANE=21477783; PubMed=11594591;
MIDIANE=21477783; PubMed=11594591;
MIDIANE=21477783; PubMed=11594591;
MIDIANE=21477783; PubMed=11594591;
MIDIANE=21477783; PubMed=11594591;
MIDIANE=2147783; PubMed=11594591;
MIDIANE=21477891591;
MIDIANE=21477891;
MIDIANE=21477891;
MIDIANE=21477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation updat
DNA 99rase B subunit (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tenacibaculum ovolyticum.
Bacteria; Bacteroidetes; Flavobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Name=gyrB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
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52920 MW; 1E1E631B9CEDFCC2 CRC64;
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85.7%;
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Pred. No.
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                                                                                            (ATP-hydrolyzing) activity; IEA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            474
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                                                                                                                                                                                                                                                                                                 topoisomerase family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Flavobacteriales;
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Best Local S
Matches 6
C -!- SIMILARITY: Belongs to the type II topoisomerase family.

R EMBL; AB034218; BAB12473.1; -.

R EMBL; AB034215; BAB12473.1; -.

R GO; GO:0005524; F:ATP binding; IEA.

GO; GO:000557; F:DNA binding; IEA.

GO; GO:0003594; F:NNA topoisomerase (ATP-hydrolyzing) activity;

R GO; GO:0006304; F:DNA modification; IEA.

GO; GO:0006304; F:DNA modification; IEA.

R GO; GO:0006505; P:DNA topological change; IEA.

R InterPro; IPR001559; ATPbind ATPase.

InterPro; IPR001594; ATPbind ATPase.

InterPro; IPR001241; DNA topoisoII.

InterPro; IPR001241; DNA topoisoII.

InterPro; IPR006171; TOprim dom.

R InterPro; IPR006171; TOprim dom.

R Ffam; PF00204; DNA gyraseB; 1.

R Pfam; PF00204; DNA gyraseB; 1.

R Pfam; PF001751; TOprim; 1.

R Pfam; PF001751; TOprim; 1.

R Pfam; PF001751; TOprim; 1.

R Pfam; PF00333; DNA gyrase_B; 1.

R PSNNRT; SM00433; TOP2C; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NON TER
                                                                                                                                                                                                                                                                                                                                                                                                                               Suzuki M., Nakagawa Y., Harayama S., Yamamoto S.;
"Phylogenetic analysis and taxonomic study of marine Cytophaga-like
bacteria: proposal for Tenacibaculum gen. nov. with Tenacibaculum
maritimum comb. nov. and Tenacibaculum ovolyticum comb. nov., and
description of Tenacibaculum mesophilum sp. nov. and Tenacibaculum
                                                                                                                                                                                                                                                                                                 amylolyticum sp. nov.";
Int. J. Syst. Evol. Microbiol. 51:1639-1652(2001).

-i- FUNCTION: DNA gyrase negatively supercoils closed circular doublestranded DNA in an ATP-dependent manner and also catalyzes the interconversion of other topological isomers of double-stranded DNA rings, including catenanes and knotted rings (By similarity).

-i- CATALYTIC ACTIVITY: ATP-dependent breakage, passage and rejoining of double-stranded DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  O9ETV9; PRELIMINARY;

O9ETV9; PRELIMINARY;

01-MAR-2001 (TrEMBLrel. 16,

01-MAR--2001 (TrEMBLrel. 27,

05-JUL-2004 (TrEMBLrel. 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pfam; PF00204; DNA GyraseB; 1.
Pfam; PF02518; HATFase c; 1.
Pfam; PF01751; TOPDrim; 1.
PRINTS; PR00418; TP12FAMILY.
PRODOM; PD19633; TOP2c; 1.
SMART; SM00433; TOP2c; 1.
SMART; SM00433; TOP2c; 1.
SMART; SM00177; TOP0ISOMERASE_II; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=MBIC1544, and IFO16022;
MEDLINE=21477783; PubMed=11594591;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria, Bacteroidetes, Flavobacteria, Flavobacteriales, Flavobacteriaceae, Cellulophaga.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cytophaga lytica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
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IPR011558; DNA_gyrase_B.
IPR001241; DNA_topoisoII.
IPR006171; Toprim_dom.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        474 AA;
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53026 MW;
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Last annotation update)
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Pred. No.
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Query Match
Best Local S
Matches
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                              InterPro; IPR003594; ATPbind_ATPase.
InterPro; IPR011558; DNA gyrase_B.
InterPro; IPR0012241; DNA topoisoII.
InterPro; IPR006171; Toprim_dom.
Pfam; PF00204; DNA gyraseB; 1.
Pfam; PF002518; HATPase_c; 1.
Pfam; PF01751; Toprim; 1.
PRINTS; PR00418; TP12FAMILY.
PRINTS; PR00418; TP12FAMILY.
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ATP-binding; Is,
NON_TER 1
NON_TER 474
SEQUENCE 474
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amylolyticum sp. nov.";
amylolyticum sp. nov.";
Int. J. Syst. Evol. Microbiol. 51:1639-1652(2001).

-i. FUNCTION: DNA gyrase negatively supercoils closed circular double-
istranded DNA in an ATP-dependent manner and also catalyzes the 
interconversion of other topological isomers of double-stranded 
interconversion of other topological isomers of the propological isomers of the propological isomers of the propological
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Q9F1Y8;
01-MAR-2001
01-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA rings, including catenanes and knotted rings (By sim -: CATALYTIC ACTIVITY: ATP-dependent breakage, passage and of double-stranded DNA.

-: SIMILARITY: Belongs to the type II topoisomerase family.

EMBL; AB034232; BAB12490.1; -.

EG; GO:000524; F:ATP binding; IEA.

GO; GO:0003677; F:DNA binding; IEA.

GO; GO:0003918; F:DNA topoisomerase (ATP-hydrolyzing) activical GO; GO:0006265; P:DNA topoisomerase (ATP-hydrolyzing) activical GO; GO:0006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-21477783; PubMed=11594591; Suzuki M., Nakagawa Y., Harayama S.; Yamamoto S.; Suzuki M., Nakagawa Y., Harayama S.; Yamamoto S.; Phylogenetic analysis and taxonomic study of marine Cytophaga-like bacteria: proposal for Tenacibaculum gen. nov. with Tenacibaculum maritimum comb. nov. and Tenacibaculum ovolyticum comb. nov., and description of Tenacibaculum mesophilum sp. nov. and Tenacibaculum
                                                                                                                                                                                                                       ProDom; PD149633; DNA_gyrase_B; 1.
SMART; SM00433; TOP2C; 1.
PROSITE; PS00177; TOPDISCMERASE_II; 1.
ATP-binding; Isomerase; Topoisomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tenacibaculum ovolyticum.
Bacteria; Bacteroidetes; Flavobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
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6; Conserv
6; Conserv
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                                                                                                                             474
474 AA;
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474 AA;
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(TrEMBLrel.
      Conservative
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                                                                                                                             53068 MW; 7FEFA02AE6E986F9 CRC64;
                                 89.2%;
85.7%;
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16,
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Pred. No.
                                    Score 33;
Pred. No.
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      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ATP-hydrolyzing) activity;
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                                                                     DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB
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                                                                  Length 474;
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YSSNLHS

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164

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Best Local S
Matches 6
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Q9F1Z0;
01-MAR-2001
01-MAR-2001
01-MAR-2004
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                                                                                                                                                                                                                                   PROSITE;
ATP-bindi
                                                                                                                                                                                                                                                                                                                                                                                                                                                               amylolyvicum sp. nov.";
Int. J. Syst. Evol. Microbiol. 51:1639-1652(2001).
-!- FUNCTION: DNA gyrase negatively supercoils closed circular double-
stranded DNA in an ATP-dependent manner and also catalyzes the 
interconversion of other topological isomers of double-stranded 
DNA rings, including catenanes and knotted rings (By similarity).
-!- CATALYTIC ACTIVITY: ATP-dependent breakage, passage and rejoining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Suzuki M., Nakagawa Y., Harayama S., Yamamoto S:;
"Phylogenetic analysis and taxonomic study of marine Cytophaga-like
bacteria: proposal for Tenacibaculum gen. nov. with Tenacibaculum
maritimum comb. nov. and Tenacibaculum ovolyticum comb. nov., and
description of Tenacibaculum mesophilum sp. nov. and Tenacibaculum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         O9F1Y9 PRELIMINARY; PRT; 474 AA.
Q9F1Y9;
O1-MAR-2001 (TrEMBLrel. 16, Created)
O1-MAR-2001 (TrEMBLrel. 16, Last sequence update)
O1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
                                                                                                                                                                                                                                          ProDom; PD149633; DNA gyrase_B; 1.
SMART; SM00433; TOP2c; 1.
PROSITE; PS00177; TOPOISOMERASE_II; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tenacibaculum maritimum.
Bacteria; Bacteroidetes; Flavobacteria; Flavobacteriales;
Flavobacteriaceae; Tenacibaculum.
                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=21477783;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Name=gyrB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA gyrase B subunit (Fragment).
                                                                                                      164
                                                                                                                                                                                                                                inding; Isomerase; Topoisomerase.
                                                                                                                            μ.
                                                                                                                                                  Similarity 6; Conserv
                                                                                                                            YSSNLHS 7
                                                                                                      YSENLHS 170
(TrEMBLrel. 16,
(TrEMBLrel. 16,
(TrEMBLrel. 26,
                                                                                                                                                                                                474 AA;
                                                                                                                                                                                                            474
                                                                                                                                                    Conservative
                                               PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PubMed=11594591;
                                                                                                                                                                                              52888 MW;
                                                                                                                                                             89.2%;
                      Created)
                                                                                                                                                  <u>.</u>
Last sequence update)
Last annotation update)
                                                                                                                                                           Score 33; DB 2;
Pred. No. 1.5e+02;
                                                                                                                                                                                                47D1F2DBB02B996E CRC64;
                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                              II topoisomerase family
                                                                                                                                                                                                                                                                                                                                                                                                (ATP-hydrolyzing) activity; IEA
                                               474
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amylolyticum Sp. nov.";

CC -: FUNCTION: DNA gyrase negatively supercoils closed circular double-
cc stranded DNA in an ATP-dependent manner and also catalyzes the
interconversion of other topological isomers of double-stranded
CC DNA rings, including catenanes and knotted rings (By similarity).
CC -: CATALYTIC ACTIVITY: ATP-dependent breakege, passage and rejoining
CC -: SIMILARITY: Belongs to the type II topoisomerase family.
CC -: SIMILARITY: Belongs to the type II topoisomerase family.
CC -: SIMILARITY: Belongs to the type II topoisomerase family.
CR MBL; AB034228; BAB12486.1; -.

DR GO; GO:0003577; F:DNA topoisomerase (ATP-hydrolyzing) activity; IEA.
CG; GO:0003578; F:DNA topoisomerase (ATP-hydrolyzing) activity; IEA.
CG; GO:0003518; F:DNA topoisomerase (ATP-hydrolyzing) activity; IEA.
CG; GO:0003625; P:DNA topoisomerase (ATP-hydrolyzing) activity; IEA.
CG; GO:0006265; P:DNA topoisomerase (ATP-hydrolyzing) activity; IEA.
CG; GO:0006304; P:DNA topoisomerase (ATP-hydrolyzing) activity; IEA.
CG; GO:0006265; P:DNA topoisomerase (ATP-hy
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Best Local Similarity
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NON_TER
SEQUENCE
                                                                                                                                                                                                                                      01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
                    SEQUENCE FROM N.A.
STRAIN=DSM5423;
MEDLINE=21477783; PubMed=11594591;
                                                                                                                    Psychroflexus gondwanensis.
Bacteria; Bacteroidetes; Flavobacteria; Flavobacteriales; Flavobacteriaceae; Psychroflexus.
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  Suzuki M., Nakagawa
                                                                                                                                                                                                                    DNA gyrase B subunit (Fragment).
                                                                                                                                                                                                                                                                                                           Q9FAV1;
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MEDLINE=21477783; PubMed=11594591;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PROSITE; PS00177; TOPOISOMERASE II; 1.
ATP-binding; Isomerase; Topoisomerase.
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                                                                                                             _TaxID=251;
                                                                                                                                                                                                                                                                                                                                                                                                                                             164
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      474 AA;
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                                                                                                                                                                                                                                                                                                                               PRELIMINARY;
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  Harayama
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Pred. No. 1.5e+02
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    Yamamoto
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Best Local
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InterPro; IPR01158; DNA gyrase B.
InterPro; IPR011241; DNA topoisoII.
InterPro; IPR001241; Toprim dom.
Pfam; PF00204; DNA gyraseB; 1.
Pfam; PF002518; HATPase c; 1.
Pfam; PF01751; Toprim; 1.
PRINTS; PR00418; TP12FAMILY.
PRODOM; PD149633; DNA gyrase B; 1.
                                                  Suzuki M., Nakagawa Y., Harayama S., Yamamoto S.;
"Phylogenetic analysis and taxonomic study of marine Cytophaga-like
bacteria: proposal for Tenacibaculum gen. nov. with Tenacibaculum
maritimum comb. nov. and Tenacibaculum gen. nov. and Tenacibaculum
description of Tenacibaculum mesophilum sp. nov. and Tenacibaculum
amylolyticum sp. nov.";
Int. J. Syst. Evol. Microbiol. 51:1639-1652(2001).
-i- FUNCTION: DNA gyrase negatively supercoils closed circular doubl
stranded DNA in an ATP-dependent manner and also catalyzes the
interconversion of other topological isomers of double-stranded
interconversion of other topological isomers of double-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cyrophaga lytica.
Bacteria; Bacteroidetes; Flavoba
Bacteria; Cellulophaga.
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Q9FAV9;
01-MAR-2001 (TrEMBLrel. 16,
01-MAR-2001 (TrEMBLrel. 26,
01-MAR-2004 (TrEMBLrel. 26,
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SEQUENCE
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-!- CATALYTIC ACTIVITY: ATP-dependent breakage, passage and rejoining of double-stranded DNA.

-!- SIMILARITY: Belongs to the type II topoisomerase family.

EMBL; AB034226; BAB12484.1; -.

GO; GO:0005524; F:ATP binding; IEA.

GO; GO:0003518; F:DNA binding; IEA.

GO; GO:0003318; F:DNA topoisomerase (ATP-hydrolyzing) activity; IEA.

GO; GO:0006304; P:NA modification; IEA.

GO; GO:0006305; P:DNA modification; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A. STRAIN=IFO15986;
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                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=21477783; PubMed=11594591;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Name=gyrB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA gyrase B
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ATP-binding; Isomerase; Topoisomerase.
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FUNCTION: DNA gyrase negatively supercoils closed circular double-
      DNA rings, including catenanes and knotted rings (By similarity) CATALYTIC ACTIVITY: ATP-dependent breakage, passage and rejoinin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            164
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Last annotation update)
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Pred. No. 1.5e+02;
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                                                                                                                     closed circular double-
passage and
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RESULT

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Best Local S
Matches 6
                                                                                                   amylolyticum sp. nov.";

Lint. J. Syst. Evol. Microbiol. 51:1639-1652(2001).

Lint. J. Syst. Evol. Microbiol. 51:1639-1652(2001).

C. -!- FUNCTION: DNA gyrase negatively supercoils closed circular double-stranded DNA in an ATP-dependent manner and also catalyzes the interconversion of other topological isomers of double-stranded C DNA rings, including catenanes and knotted rings (By similarity).

C. -!- CATTALYTIC ACTIVITY: ATP-dependent breakage, passage and rejoining of double-stranded DNA.

C. -!- SIMILARITY: Belongs to the type II topoisomerase family.

REMBL; AB034216; BAB12474.1;

CR GO: GO:000357; F:ATP binding; IEA.

GO; GO:0003573; F:DNA binding; IEA.

GO; GO:0003573; F:DNA binding; IEA.

GO; GO:0003574; P:DNA modification; IEA.

GO; GO:0003575; P:DNA modification; IEA.

GO; GO:0006365; P:DNA topoisomerase (ATP-hydrolyzing) activity; IEA.

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G0; G0:0003677; F:DNA binding; IEA.
G0; G0:0003187; F:DNA topoisomerase (A:
G0; G0:000318; F:DNA topoisomerase (A:
G0; G0:0006304; P:DNA modification; IEE
G0; G0:0006265; P:DNA topological chang
InterPro; IPR001594; ATPbind_ATPase.
InterPro; IPR001595; DNA gyrase B.
InterPro; IPR001241; DNA topoisoII.
InterPro; IPR001241; TOprim dom.
Pfam; PF00204; DNA gyraseB; 1.
Pfam; PF00218; HATPase_c; 1.
Pfam; PF00218; HATPase_c; 1.
Pfam; PF01518; TOprim; 1.
Pfam; PF0149633; DNA_gyrase_B; 1.
SNART; SN00433; TOP2C; 1.
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i-SIMILARITY: Belongs to the type II topoisc
EMBL; AB034217; BAB12475.1;
GO; GO:0005524; F:ATP binding; IEA.
GO; GO:000577; F:DNA binding; IEA.
GO; GO:0003677; F:DNA binding; IEA.
GO; GO:0006304; F:DNA modification; IEA.
GO; GO:0006304; P:DNA topological change; IEA.
GO; GO:0006265; P:DNA topological change; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Suzuki M., Nakagawa Y., Harayama S., Yamamoto S.; "Phylogenetic analysis and taxonomic study of marine Cytophaga-like bacteria: proposal for Tenacibaculum gen. nov. with Tenacibaculum marithmum comb. nov. and Tenacibaculum ovolyticum comb. nov., and description of Tenacibaculum mesophilum sp. nov. and Tenacibaculum
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01-MAR-2001 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
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STRAIN=IAM14306
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q9FAW0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=21477783; PubMed=11594591;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Bacteroidetes; Flavobacteria; Flavobacteriales;
Flavobacteriaceae; Cellulophaga.
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ATP-binding; Isomerase; Topoisomerase.
                                                                            InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                164
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IPR003594; ATPbind_ATPase.
IPR011558; DNA_gyrase_B.
IPR001241; DNA_topoisoII.
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26,
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Pred. No.
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RESULT 20
Q9FAW1
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    TTERREARED COCCEPTITION ARE RESERVED COCCEPTION OF THE RESERVED COCCEPTION 
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Best Local Similarity
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NON_TER
SEQUENCE
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Pfam; PF02518; HATPase c; 1.
Pfam; PF01751; Toprim; 1.
PRINTS; PR00418; TPI2FAMILY.
ProDom; PD149633; DNA_gyrase_B; 1.
SMART; SM00433; TOP2c; 1.
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MEDLINE=21477783; PubMed=11594591;
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Bacteria; Bacteroidetes; Flavobacteria; Flavobacteriales;
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
DNA gyrase B subunit (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Phylogenetic analysis and taxonomic study of marine Cytophaga-like bacteria: proposal for Tenacibaculum gen. nov. with Tenacibaculum maritimum comb. nov. and Tenacibaculum ovolyticum comb. nov., and description of Tenacibaculum mesophilum sp. nov. and Tenacibaculum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Suzuki M., Nakagawa Y., Harayama S., Yamamoto S.;
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PROSITE; P800177; TOPOISOMERASE_II; 1.
ATP-binding; Isomerase; Topoisomerase.
NON_TER 1 1
NON_TER 474 474
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85.7%;
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Pred. No. 1.5e+02;
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                                                                                                                                        Query Match
Best Local S
Matches 6
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C. -- FUNCTION: DNA gyrase negatively supercoils closed circular double-
ce interanded DNA in an ATP-dependent manner and also catalyzes the 
interconversion of other topological isomers of double-stranded 
DNA rings, including catenanes and knotted rings (By similarity). 
C. --- CATALYTIC ACTIVITY: ATP-dependent breakage, passage and rejoining 
C. --- SIMILARITY: Belongs to the type II topoisomerase family. 
EMBL; AB034213; BAB12471.1; --- 
NR GO; GO:0003524; F:ATP binding; IEA. 
GO; GO:0003577; F:DNA binding; IEA. 
GO; GO:0003677; F:DNA topoisomerase (ATP-hydrolyzing) activity; IEA. 
RGO; GO:000364; P:DNA topoisomerase (ATP-hydrolyzing) activity; IEA. 
RGO; GO:0006265; P:DNA topoisomerase; IEA. 
RGO; GO:000
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Best Local
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InterPro; IPR001241; DNA_topoisoII.
InterPro; IPR001241; DNA_topoisoII.
InterPro; IPR006171; Toprim_dom.
Pfam; PF00204; DNA_gyraseB; 1.
Pfam; PF02518; HATPase_c; 1.
Pfam; PF01751; Toprim; 1.
PRINTS; PR00418; TPIZFAMILY.
PRODOm; PD149633; DNA_gyrase_B; 1.
SMART; SM0433; TOP2C; 1.
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                                                                                                                                                                                                                                                                                                                     NON_TER
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01-MAR-2001
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ATP-binding; Isomerase; Topoisomerase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=21477783; PubMed=11594591;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21
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   164
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YSENLHS 170
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                                                                                                                                                                                                                                                                                    474 AA;
                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                        53119 MW;
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85.7%;
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16, Last sequence update)
26, Last annotation update)
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                                                                                                                                    Score 33; DB 2; 1
Pred. No. 1.5e+02;
0; Mismatches 1
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Pred. No.
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                                                                                                                                                                                                                                                                                        38D059D7E0FE1F93 CRC64;
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PERMY 22
QPFAWA
ID QPFAWA
ID QPFAWA
AC QPFAWA

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Q7PX57
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RI Int. J. Syst. Evol. Microbiol. 51:639-1652(2001).

CC -!- FUNCTION: DNA gyrase negatively supercoils closed circular double-
CC stranded DNA in an ATP-dependent manner and also catalyzes the
interconversion of other topological isomers of double-stranded
CC DNA rings, including catenanes and knotted rings (By similarity).

CC -!- CATALYTIC ACTIVITY: ATP-dependent breakage, passage and rejoining
CC of double-stranded DNA.

CC -!- SIMILARITY: Belongs to the type II topoisomerase family.

DR GO; GO:0005524; F:ATP binding; IEA.

CR GO; GO:0005527; F:DNA binding; IEA.

CG; GO:0003577; F:DNA binding; IEA.

CG; GO:000357; F:DNA modification; IEA.

CG; GO:0003504; P:DNA modification; IEA.

CG; GO:0006504; P:DNA modification; IEA.

CG; GO:0006504; P:DNA modification; IEA.

CG; GO:0006507; F:DNA topological change; IEA.

CR GO; GO:0006517; Topxin_dom.

CR InterPro; IPR001241; DNA_TopoisoII.

DR InterPro; IPR001241; DNA_TopoisoII.

DR Ffam; PF00204; DNA gyraseB; 1.

CR Ffam; PF00204; DNA gyraseB; 1.

CR Ffam; PF00204; DNA_ByraseB; 1.

DR Pfam; PF001751; Toprin; 1.

DR Pfam; PF00177; TOPOISOMERASE II; 1.

DR PGDOIS ; PF00177; TOPOISOMERASE II; 1.
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Best Local S
Matches 6
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Q9FAW4;
01-MAR-2001
01-MAR-2001
01-MAR-2004
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NON TER
SEQUENCE
                             O7PX57; PRELIMINARY; PRT; 817 AA.
07PX57; O1-MAR-2004 (TrEMBLrel. 26, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence up
01-MAR-2004 (TrEMBLrel. 26, Last annotation of the company of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Suzuki M., Nakagawa Y., Harayama S., Yamamoto S.; "Phylogenetic analysis and taxonomic study of marine Cytophaga-like bacteria: proposal for Tenacibaculum gen. nov. with Tenacibaculum maritimum comb. nov. and Tenacibaculum ovolyticum comb. nov., and description of Tenacibaculum mesophilum sp. nov. and Tenacibaculum mesophilum sp. nov.
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Bacteria; Bacteroidetes; Flavobacteria; Flavobacteriales;
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation updat
DNA 9yrase B subunit (Fragment).
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480 i
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53889 MW;
Arthropoda;
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85.7%;
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Pred. No.
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Hexapoda; Insecta; Pterygota;
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annotation update)
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Best Local S
Matches 6
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del Campillo E., Bennett A.B.;
"pedicel breakstrength and cellulase gene expedicel breakstrength and cellulase gene expedicel breakstrength and cellulase gene expedicate abscission.";
plant Physiol. 111:813-820(1996).
GO; GO:0004553; F:hydrolase activity, hydroly GO; GO:0005975; P:carbohydrate metabolism; IE InterPro; IPR001701; Glyco_hydro_9.
InterPro; IPR008928; Glyco_trans_6hp.
Pfam; PF00759; Glyco_hydro_9; 1.
NON TER
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Q9S950;
01-MAY-2000 (TrEMBLrel. 13, C
01-MAY-2000 (TrEMBLrel. 13, L
01-OCT-2003 (TrEMBLrel. 25, L
CEL2=CELLULASE 2 (Fragment).
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NON_TER
SEQUENCE
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NON TER
SEQUENCE
                                                                                     05-JUL-2004 (TrEMBLrel. 27, Create 05-JUL-2004 (TrEMBLrel. 27, Last s 05-JUL-2004 (TrEMBLrel. 27, Last pottative endo-1,4,-beta-glucanase
                                                                                                            Q70Y33;
05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
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Submitted (MAR-2002) to the EMBL/GenBank/DDBJ
-i- CAUTION; The sequence shown here is derive
EMBL/GenBank/DDBJ whole genome shotgun (WG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Neoptera; Endopter NCBI_TaxID=180454;
Lycopersicon esculentum (Tomato).
Eukaryota; Viridiplantae; Strepto
Spermatophyta; Magnoliophyta; eud
                                                                                                                                                                                                    Q70Y33
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; aster
lamiids; Solanales; Solanaceae; Solanum.
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                                                                   Name=cel;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              preliminary data.
; AAAB01008987; EAA01793.1; -.
                                                                                                                                                                                                                                                                                                                   143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       589
                                                                                                                                                                                                                                                                                                                                                           1 YSSNLHS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                       9
                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                 |||:|||
YSSSLHS 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 YSSNLH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       YSSNLH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Endopterygota; Diptera; Nematocera; Culicoidea; Anopheles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       817 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               169 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      169
                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                    PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                               7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      169
                                                                                                                                                                                                                                                                                                                                                                                                                            86.5%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18674 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       87412 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         89.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
  Streptophyta; Embryo
/ta; eudicotyledons;
                                                                                                            Last sequence update)
Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Last sequence update)
Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Created)
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                                                                                                                                                        Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 33;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                              Score 32;
Pred. No.
                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       F898FB905B8B2995 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CE8AB3AC5637C481 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                       (Fragment).
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re is derived from an
shotgun (WGS) entry which
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hydrolyzing O-glycosyl
                                                                                                                                                                                                                                                                                                                                                                                                                              78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 2; Lo
2.8e+02;
  Embryophyta;
edons; core e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Length 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IEA.
                                                                                                               update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 817;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                           Indels
    a; Tracheophyta;
eudicots; aster
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                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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RESULT 27
Q4375
ID Q4375
AC Q4375
AC Q4375
AC Q4375
DT 01-NC
DT 01-NC
DT 01-QC
DT 01
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Best Local S
Matches 6
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Best Local Similarity 85.
Matches 6; Conservative
                                                                                                                                                 Q43751 PRELIMINARY;
Q43751;
Q43751;
Q1-00V-1996 (TrEMBLrel. 01, C:
Q1-00V-1996 (TrEMBLrel. 25, Li
Q1-0CT-2003 (TrEMBLrel. 25, Li
Cellulase (EC 3.2.1.4) (Fragma
Name=ccx3;
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CAD44274;
02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
Putative endo-1,4,-beta-glucanase (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NON_TER
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Submitted (AUG-2002) to the EMBL/GenBank/DDBJ
EMBL; AJ505749; CAD44274.1; -
InterPro; IPR001701; Glyco_hydro_9.
InterPro; IPR008928; Glyco_trans_6hp.
Pfam; PF00759; Glyco_hydro_9; 1.
Capsicum annuum (Bell pepper).
Bukaryota; Viridiplantae; Streptophyta;
Spermatophyta; Magnoliphyta; eudicotyle
lamiids; Solanales; Solanaceae; Capsicum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledona; core eudicots; Asteridae; lamiida; Solanalea; Solanaceae; Solanum; Lycopersicon.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (AUG-2002) to the EMBL; AJ505749; CAD44274.1; NON_TER 1 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=cv. Arka vikas, TISSUE=Fruit;
Saiprasad G.V.S., Lalitha A.;
"Isolation, cloning and characterization of endo-1,4,-beta-glucanase
gene from ripening tomato fruit (Lycopersicum esculentum cv. Arka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=4081;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lycopersicon esculentum (Tomato).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Fruit;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            144
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            YSSSLHS 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 YSSNLHS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                YSSNLHS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  172 172
172 AA; 19004 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               172 172
172 AA; 19004 MW; F8584877ABEB83F9 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Solanaceae; Solanum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              86.5%;
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                                                                                                                                                                                                                                 Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                              Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 32; DB
Pred. No. 80;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 32; DB 2;
Pred. No. 80;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      F8584877ABEB83F9 CRC64;
        Capsicum.
                                                                                                                                                                                                                                                                                                                                                                                               325
                                        Embryophyta; Tracheophyta;
edons; core eudicots; aster
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 172;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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RESULT 28
Q22124
ID Q2212
AC Q2212
DT 01-VIC
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Query Match
Best Local S
Matches 6
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Best Local S
Matches 6
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Q22124;
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PIR; S61447; S61447.

GO; GO:0008810; F:cellulase activity; IEA.

GO; GO:0005975; P:carbohydrate metabolism;

InterPro; IPR001701; Glyco_hydro_9.

InterPro; IPR008928; Glyco_trans_6hp.

Pfam; PF00759; Glyco_hydro_9; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Caemorhabditis elegans.
Caemorhabditida; Rhabditoidea;
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea;
Rhabditidae; Peloderinae; Caenorhabditis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Glycosidase; Hydrolase.

NON_TER 1 1

NON_TER 325 325
                                                                                     EMBL; U41272; AAA82450.2; -. PIR; T29924; T29924. InterPro; IPR003839; DUF215. Pfam; PF02688; DUF215; 1. SEQUENCE 331 AA; 37829 MW;
                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
STRAIN=Bristol N2;
Geisel C., Gattung S.;
Submitted (DEC-1995) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-NOV-1996 (TrEMBLrel.
01-JUN-2003 (TrEMBLrel.
01-OCT-2003 (TrEMBLrel.
                                                                                                                                                                                                       Waterston R.;
Submitted (JAN-2003) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                               "Genome sequence of the nematode C. elegans: a platform for investigating biology. The C. elegans Sequencing Consortium Science 282:2012-2018(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A. STRAIN=Bristol N2; MEDLINE=99069613;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=4072;
                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
STRAIN=Bristol N2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wilson R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RIBDALTIDES; Peloderinae; NCBI_TaxID=6239;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mol. Biol.
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-JUN-2003 (TrEMBLrel. 24,
-OCT-2003 (TrEMBLrel. 25,
elegans SRU-48 protein.
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Similarity 6; Conserv
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325 AA;
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36087 MW;
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85.7%;
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Pred. No.
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Pred. No. 1.6e+02;
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                                                                                          7FC40A485BB9F837 CRC64;
Mismatches
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                          1.7e+02
                                             DB 2;
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CC -!- SIMILARITY: Belongs to class-II aminoacyl-tRNA synthetase family.

DR EMBL; AP004594; BAC12522.1; -.

DR GO; GO:0005737; C:cytoplasm; IEA.

GO; GO:0005524; F:ATP binding; IEA.

GO; GO:0005524; F:ATP binding; IEA.

GO; GO:0005524; F:Proline-tRNA injase activity; IEA.

GO; GO:0004527; F:proline-tRNA injase activity; IEA.

GO; GO:000643; P:prolyl-tRNA aminoacylation; IEA.

GO; GO:000643; P:prolyl-tRNA aminoacylation; IEA.

GO; GO:0006412; P:protein biosynthesis; IEA.

R GO; GO:0006412; P:protein biosynthesis; IEA.

OR InterPro; IPR004194; HGTP_anticodon.

InterPro; IPR004194; ProS_fam_I.

DR InterPro; IPR002316; tRNA-synt_pro.

R InterPro; IPR002316; tRNA-synt_pro.

DR InterPro; IPR006195; tRNA-synt_pro.

DR Ffam; PF03129; HGTP_anticodon; 1.

DR Pfam; PF03129; HGTP_anticodon; 1.
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Matches
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Q6DBD4;
Q1-OCT-2004
01-OCT-2004
01-OCT-2004
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Q8ESQ5;
01-MAR-2003
01-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ridge and its unexpected adaptive capabilities to extreme environments.";
Cheuk R., Chen H., Kim C.J., Shinn P., Ecker J.R.;
Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; BT015088; AB771J960.1; -
SEQUENCE 480 AA; 53687 MW; 4984BFD029021C5A CRC64;
                                                       SEQUENCE FROM N.A.
                                                                                                      Arabidopsis thaliana (Mouse-ear cress).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Aggonliophyta; eudicotyledons; core eudicots; rosids; eurosids [I] Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                         PRINTS; PR01046; TRNAŠYNTHPRO.
TIGRPAMs; TIGR00408; pro5 fam I; 1.
PROSITE; PS50862; AA TRNA_LIGĀSE II; 1.
Aminoacyl-trna synthetase; Complete proteome.
SEQUENCE 478 AA; 54750 MW; 5E37D3553928D880 CRC64;
                                                                                                                                                                   At5g16410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=22220767; PubMed=12235376; Takami H., Takaki Y., Uchiyama I.; "Genome sequence of Oceanobacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oceanobacillus iheyensis.
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                                                                                      NCBI_TaxID=3702;
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Pred. No. 2.5e.
2; Mismatches
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01-MAR-2001
01-MAR-2001
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Q96547;
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Similarity to N-hydroxycinnamoyl/benzoyltransferase.
Arabidopsis thaliana (Mouse-ear cress)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; endicotyledons; core eudicots; rosids;
                                                                                                                                                                                                                                                                                                                                                                                                                         01-FEB-1997 (TrEMBLrel. 02,
01-FEB-1997 (TrEMBLrel. 02,
01-MAR-2004 (TrEMBLrel. 26,
                                                                                                                                 TISSUE=Abscission zone;
Trainotti L., Perrarese L., Casadoro G.;
"Different endo-beta-1,4-glucanases are expressed during abscission
                                                                                                                                                                                                                                                                                   Capsicum annum (Bell pepper).

Eukaryota; Viridiplantae; Streptophyta; Embryophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eu
lamiids; Solanales; Solanaceae; Capsicum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Miyajima N., Tabata S.;
"Structural analysis of Arabidopsis thaliana
features of the 1.6 Mb regions covered by two
                                                                                                                                                                                                                                                                                                                                                                          Name=ccel3;
                                                                                                                                                                                                                                                                                                                                                                                                     Cellulase (Endo-beta-1,4-glucanase)
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GO; GO:0016740; F:transferase activity;
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                          Kluwer Academic Publishers,
[2]
                                                                         BIOLOGY AND
                                                                                                                                                                                                                 SEQUENCE
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Sato S., Kotani H., Nakamura Y., Kaneko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              eurosids II; Brassicales; Brassicaceae; Arabidopsis.
       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nterPro;
                                                fruit ripening in pepper and peach plants.";
) Kanellis A.K., Chang C., Kende and Grierson D.
LOGY AND BIOTECHOLOGY OF THE PLANT HORMONE ETHY.
wer Academic Publishers, Dordrecht (1998).
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.ucanase) (EC 3.2.1.4).
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1; Mismatches
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                                                                         ETHYLENE,
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E
                                                                                                                                                                                                                                                                                                                                      Tracheophyta;
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                                                                      pp.191-196,

    Sequence

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=Castlemart; TISSUE=Pericarp;
Brummell D.A., Lashbrook C.C., Bennet
"Plant endo-1,4-beta-D-glucanases: st
physiological function.";
ACS Symp. Ser. 566:100-129(1994).
EMBL; U1305; RAA69909.1;
PIR; T06350; T06350.
HSSP; 077044; IKSB.
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HSSP; O77044; 1KSB.

GO; GO:0008810; F:cellulase activity; IEA.

GO; GO:0005975; P:carbohydrate metabolism; IEA.

InterPro; IPR001701; Glyco_hydro_9.

InterPro; IPR008928; Glyco_trans_6hp.

Pfam; PP00759; Glyco_hydro_9; 1.

PROSITE; PS00698; GIYCOSYL_HYDROL_F9_2; UNKNOWN_1.
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01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Endo-1.4-beta-glucanase precursor (EC 3.2.1.4).
                                                                                                                                                                                                                                                                                                                                                                                                      GO; GO:0008810; F:cellulase activity; IEA.
GO; GO:0005975; P:cearbohydrate metabolism;
InterPro; IPR001701; Glyco_hydro_9.
InterPro; IPR008928; Glyco_trans_6hp.
Pfam; PF00759; Glyco_hydro_9; 1.
PROSITE; PS00698; GLYCOSYL_HYDROL_F9_2; UNI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=95086382; PubMed=7994180; Lashbrook C.C., Gonzalez-Bosch C., Bennett A.B.; "Two divergent endo-beta-1,4-glucanase genes exhibit overlapping expression in ripening fruit and abscising flowers."; Plant Cell 6:1485-1493(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=Castlemart; TISSUE=Pericarp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Viridiplantae; Streptophyta; Embryo
Spermatophyta; Magnoliophyta; eudicotyledons;
lamilds; Solanales; Solanaceae; Solanum.
                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                   Glycosidase; Hydrolase; Signal.
SIGNAL 1 23 Po
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Name=Cel2;
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                                                                                                                                   Local Similarity
les 6; Conserv
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216
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                                                                 YSSNLHS 7
YSSSLHS 222
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                                                                                                                                                                                                                                                                             489 AA;
                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                  489
                                                                                                                                                                                                                                                                                 54118 MW;
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85.7%;
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85.7%;
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                                                                                                                                                                       Score 32; I
Pred. No. 2
                                                                                                                                                                                                                                                                             Potential.
endo-1,4-beta-glucanase.
; FFADA8C4675F685F CRC64;
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Pred. No.
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                                                                                                                                          Mismatches
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edons; core eudicots; aster
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                                                                                                                                          Indels
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Q93WZ0
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STRAIN-ARCC 49582;
STRAIN-ARCC 49582;
STRAIN-ARCC 49582;
Submitted (MAR-2003) to the EMBL/GenBank/DDBJ databases.
-I- SIMILARITY: Belongs to the type II topoisomerase fam
EMBL; AY257486; AAP13529.1;
-GO; GO:0005524; F:ATP binding; IBA.
GO; GO:0003677; F:DNA binding; IBA.
GO; GO:0003918; F:DNA binding; IBA.
GO; GO:000318; F:DNA modification; IBA.
GO; GO:0006304; P:DNA modification; IBA.
GO; GO:0006305; P:DNA modification; IBA.
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Q83Z58;
01-JUN-2003
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; aster
lamiids; Solanales; Solanaceae; Nicotiana.
NCBI_TaxID=4097;
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01-DEC-2001 (TYEMBLrel. 19, Last sequence update)
01-MAR-2004 (TYEMBLrel. 26, Last annotation update)
Endo-beta-1,4-glucanase precursor (EC 3.2.1.4).
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01-DEC-2001
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01-OCT-2003
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MEDLINE=21480065; PubMed=11595799;
Goellner M., Wang X., Davis E.L.;
"Endo-beta-1,4-glucanase expression"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q93WZ0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Name=gyrB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA gyrase subunit
                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=228419;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Entomoplasmataceae;
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Bacteria; Firmicutes; Mollicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nteractions."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 85.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                YSSSLHS
       IPR003594;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 36
500 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pleciae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (TrEMBLrel.) (TrEMBLrel.) (TrEMBLrel.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mesoplasma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     55075 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           86.5%;
85.7%;
       ATPbind
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Fragment).
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24, Last sequence update)
25, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 32;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Potential.
endo-beta-1,4-glucanase.
; 28BB48D421165798 CRC64;
       ATPase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Entomoplasmatales;
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                                                                                                      (ATP-hydrolyzing) activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           536
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                                                                                                             IEA.
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RESULT 37
Q9RP84
ID Q9RP8
AC Q9RP8
DT 01-M2
DT 01-M2
DT 01-M2
DT 01-Q2
DT 0
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Best Local S
Matches 6
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Best Local S
Matches 6
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InterPro; IPR001241; DNA topoisoII.
InterPro; IPR001271; Toprim dom.
Pfam; PP00204; DNA gyraseB; 1.
Pfam; PP002518; HATPase_C; 1.
Pfam; PP01751; Toprim; 1.
Pfam; PP01751; Toprim; 1.
PRINTS; PR00418; TP12FAMILY.
ProDom; PD149633; DNA gyrase B; 1.
SMART; SM00387; HATPase_C; 1.
SMART; SM00433; TOP2C; 1.
SMART; SM00433; TOP2C; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q7Q8M6
Q7Q8M6;
Q7Q8M6;
01-MAR-2004 (TrEMBLrel. 2
01-MAR-2004 (TrEMBLrel. 2
01-MAR-2004 (TrEMBLrel. 2
                                                                                                 Q9RP84;
Q9RP84;
01-MAY-2000
01-MAY-2000
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anopheles Genome Sequencing Consortium;
Submitted (MAR-2002) to the EMBL/GenBank/DDBJ databases.
-!- CAUTION: The sequence shown here is derived from an EMBL/GenBank/DDBJ whole genome shotgun (WGS) entry v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anopheles gambiae str. PEST.
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Nematocera; Culicoidea; Anopheles.
   Name=gyrB;
Acholeplasma laidlawii
                                                                    DNA gyrase beta subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pfam; P
NON_TER
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Name=agCG49789; ORFN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=PEST;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=180454;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROSITE; PS00177; TOPOISOMERASE_II; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SL; AAAB011008944; EAA10074.1; -.
GO:0042755; C:GPI-anchor transamidase complex; IEA.
GO:0016021; C:integral to membrane; IEA.
erPro; IPR007245; Gp116.
m; PF04113; Gp116; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             preliminary data.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          YSSNLHS 7
                                                                                                                                                                                                                                                                                                                                                                                                                         SHYNSSY
                                                                                                 (TrEMBLrel. 13, (TrEMBLrel. 13, (TrEMBLrel. 25,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   598 AA; 67546 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           536 AA; 60040 MW;
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                                                                                                                                                                                                                                             PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORFNames=ENSANGG00000018304;
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85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           86.5%;
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Last annotation update)
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                                                                                                 Last sequence update)
Last annotation update)
                                                                                                                                                                        Created)
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Pred. No. 3.2e+02;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   260B7B630D02C13A CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 598;
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Best Local S
Matches 6
                                                                           OBRFN2;
01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last seg
01-OCT-2003 (TrEMBLrel. 25, Last ann
ABC transporter ATP-binding protein.
OrderedLocusNames=FN0660;
 SEQUENCE FROM N.A.
STRAINSATCC 25586;
MEDLINE=21886394; PubMed=11889109;
Kapatral V., Anderson I., Ivanova |
Bhattacharyya A., Bartman A., Gard
                                                                                                                       Q8RFN2
                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Analysis of genes, coding for DNA g
Acholeplasma laidlawii PG-8B.";
Mol. Biol. (Mosk.) 34:292-299(2000).
                                                             Bacteria; Fusobacteria;
                                                                     Fusobacterium nucleatum
                                                                                                                                                                                                                             ATP-binding;
                                                                                                                                                                                                                                     TIGRFAMS; TIGR01059; gyrB; 1.
PROSITE; PS00177; TOPOISOMERASE_II; 1.
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Abramycheva N.Iu, Govorun V.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=20242229; PubMed=10779957;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
STRAIN=PG-8B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=2148;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Firmicutes; Mollicutes; Acholeplasmatales; Acholeplasmataceae; Acholeplasma.
                                             CBI_TaxID=76856;
                                                                                                                                      38
                                                                                                                                                             267
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                                                                                                                                                                                         Similarity 6; Conserv
                                                                                                                                                            YSPNLHS 273
                                                                                                                                                                                                                       615
                                                                                                                                                                                         Conservative
                                                                                                                        PRELIMINARY;
                                                                                                                                                                                                                               Isomerase;
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                                                                                                                                                                                                86.5%;
85.7%;
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                                                                                                                                                                                                                      Topoisomerase.
                                                             Fusobacterales;
                                                                     (subsp. nucleatum)
   Ivanova N.,
A., Gardner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for DNA gyrase from the mycoplasma
                                                                                         Last sequence update)
                                                                                                                                                                                         0,
                                                                                                                                                                                                Score 32;
Pred. No.
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                                                                                                                        PRT;
                                                                                                                                                                                                                       4D9962C35BAB73FE CRC64;
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   Reznik G., Los T
W., Grechkin G.,
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                                                              Fusobacteriaceae;
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Best Local
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STRAIN-ATCC 43446;

MEDLINE-20127862; PubMed=10660542;

MEDLINE-20127862; PubMed=10660542;

Gilbert M., Brisson J.R., Karwaski M.F., Michniewicz J.,

Cunningham A.M., Wu Y., Young N.M., Wakarchuk W.W.;

Cunningham A.M., Wu Y., Young N.M., Wakarchuk W.W.;

Tellosynthesis of ganglioside minics in Campylobacter jejuni OH4384.

Tellosynthesis of ga
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STRAIN=ATCC 43446;
Gilbert M.;
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InterPro; IPR003439; ABC_transporter.
Pfam; PF00005; ABC_transporter; 1.
ProDom; PD00006; ABC_transporter; 1.
SMART; SM00382; AAA; 1.
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GO; GO:0005524; F:ATP binding; IEA.
GO; GO:0004009; F:ATP-binding cassette (ABC)
GO; GO:0000166; F:nucleotide binding; IEA.
GO; GO:0006810; P:transport; IEA.
PubMed=14742567;
Gilbert M., Godschalk P.C., Karwaski M.F., Ang C.W., Van Belkum A.
Li J., Wakarchuk W.W., Endtz H.P.;
"Evidence for Acquisition of the Lipooligosaccharide Biosynthesis Locus in Campylobacter jejuni GB11, a Strain Isolated from a Patie with Guillain-Barre Syndrome, by Horizontal Exchange.";
Infect. Immun. 72:1162-1165(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     J. Bacteriol. 184:2005-2018(2002).
-!- SIMILARITY: Belongs to the ABC transporter
EMBL; AE010577; AAL94856:1; -.
HSSP; Q58206; 1L2T.
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Larsen N., D'Souza M., Walu
Fonstein M., Kyrpides N.C.,
                                                                                                                                                                                                                                                                                                                                                                                             Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bacteria; Proteobacteria; Epsilonproteobacteria; Campylobacterales; Campylobacteraceae; Campylobacter.
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PROSITE; PS50893; ABC_TRANSPORTER_2; 1.
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                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
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335 AA; 37442 MW; E6ABF4100E7B4C3B CRC64;
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M., Walunas T., Pusch G., Haselkorn R.,
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Search completed: December 17, Job time: 22.9213 Becs

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Q9LAK4;
01-OCT-2000
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EMBL; AF130984; AAF31769.1; -.
EMBL; AF130984; AAF31769.1; -.
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EMBL; AF422197; AAR82873.1; -.
GO; GO:0016740; F:transferase
InterPro; IPR010446; CgtA.
Pfam; PF06306; CgtA; 1.
                                                                                                   InterPro; IPR010446; CgtA. Pfam; PF06306; CgtA; 1. Transferase.
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Campylobacteria; Epsilonproteobacteria; Campylobacterales;
Bacteria; Proteobacteria; Epsilonproteobacteria; Campylobacterales;
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NCBI_TaxID=197;
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                 GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd
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        AAB81999
AAB81998
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AAB81996
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Aab81984 Gangliosi
Abull1009 Modified
Aab81986 Gangliosi
Aab81986 Gangliosi
Aab81990 Gangliosi
Abull1013 Modified
Aar33257 Rat immun
Aar53340 KW641 H C
Aay28368 BKW641 H C
Aay28368 BKW641 HS
Aab01628 Murine im
Aab81994 Gangliosi
Aab81997 Gangliosi
Aab81997 Gangliosi
Aab81997 Gangliosi
Aab81998 Gangliosi
Aab81999 Gangliosi
Aab81998 Gangliosi
Aab81998 Gangliosi
Aab81998 Gangliosi
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Aab81989 Gangliosi
Aab81999 Gangliosi
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S

Sequence 9 AA;

Query Match
Best Local Similarity
Matches 9; Conserv

100.0%; Score 56; DB 4; ilarity 100.0%; Pred. No. 1.7e+06; Conservative 0; Mismatches 0;

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Gaps

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Length 9; Indels

co comission of the	PS Claim 4; Page 142; 183pp; Japanese.  XX  The present invention describes a mo  CC specifically with ganglioside GD3. T  CC useful in the diagnosis and therapy  CC diagnosis. The present sequence is a  CC of the invention	PT New human type complementation-determining PT and derivatives against ganglioside GD3, u PT of e.g. tumors, with low antigenicity, lit PT activity in cancer.	DR WPI; 2001-266143/27.	PI Hanai N, Shitara K, Nakamura K, Niwa YY	PA (KYOW) KYOWA HAKKO KOGYO KK.	PR 30-SEP-1999; 99JP-00278291. PR 06-APR-2000; 2000JP-00105088.	PF 29-SEP-2000; 2000WO-JP006774.	PD 05-APR-2001.	PN WO200123432-A1.	OS Mus musculus.	<pre>jlioside; GD3; complementarity ser.</pre>	ylioside GD3 specific antibody	DT 03-JUL-2001 (first entry)	AC AAB81984;	RESULT 1 AAB81984 ID AAB81984 standard; peptide; 9 AA.	ALIGNMENTS	5 40 71.4 108 4	4 40 71.4 108 3	2 40 71.4 108 2	1 40 71.4 108 2	0 40 71.4 108 2	8 40 71.4 107 8 9 40 71 4 107 8	7 40 71.4 107 8	5 40 71.4 107 8 6 40 71.4 107 8	4 40 71.4 107 4	32 40 71.4 107 2 AAK30770 33 40 71.4 107 4 AAB62087	1 40 71.4 9 3	0 40 71.4 9 3 AAB06354	8 41 73.2 113 4	7 42 75.0 107 8	6 40 75 0 107 8
The antibody which can react The antibody and its derivatives are y of tumours, particularly cancer a protein used in the exemplification	ining region-transplanted antibody D3, useful in diagnosis and therapy , little side effects but potent		wa ℝ;				•	•		determining region; CDR; antibody;	related peptide SEQ ID NO: 8.				ENTS	Aae12062 Human ant		Anti-r	Variabl	Aaw04177 Variant v	Varian	Human	Adot1565 human var Ade71454 Human ant	Conser	Aar30//0 Consensus Aab62087 Human Vl			Anti-	Ado07307	Himbr	

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HQYSKLPWT

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AABB1988
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Matches 9
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             Carreer
                       Ganglioside; GD3; complementarity determining region; CDR; antibody;
                                                    Ganglioside
                                                                               03-JUL-2001
                                                                                                         AAB81988;
                                                                                                                                 AAB81988 standard;
                                                                                                                                                                                                                                                                                                             Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 6; Page 99; 121pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Drugs containing genetically-modified antibody against ganglioside GD3, its fragment, immunocompetent cell activators or/and antitumor agents in combination, applicable in treating malignant tumor like melanoma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified ganglioside GD3 antibody associated peptide #6
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                                                                                                                                                                                                                                                        Similarity 9; Conserv
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                                                GD3 specific antibody related protein SEQ ID NO:
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Pred. No. 1.7e+06;
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RESULT 4
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Best Local
                                  WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent activity in cancer.
                                                        Hanai N,
                                                                                                     30-SEP-1999; 99JP-00278291.
06-APR-2000; 2000JP-00105088.
                                                                                                                                                                                   WO200123432-A1
                                                                                                                                                                                                          Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 22; Page 172-173; 183pp; Japanese.
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06-APR-2000; 2000JP-00105088.
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9; Conserva
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                                                       Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                  GD3 specific antibody related protein SEQ ID NO:
                                                                                                                                                                                                                                             GD3;
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                                                                                                                                                                                                                                             complementarity
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                                                                                                                                                                                                                                                                                                                                      protein; 108
                                                                                KOGYO KK.
                                                         Nakamura
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Pred. No. 0.043;
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                                                         Niwa
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                                                                                                                                                                                                                                             region; CDR; antibody;
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New human type complementation-determining region-transplanted and derivatives against ganglioside GD3, useful in diagnosis a

ed antibody and therapy

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RESULT 5
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                                                                                                           The present invention describes a monoclonal antibody which can react specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a monoclonal antibody which can react specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention
                                                                                                                                                                                                                     New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent
                                                                                                                                                                                                                                                                                                                                             30-SEP-1999; 99JP-00278291.
06-APR-2000; 2000JP-00105088.
                                                                                     Sequence 108
                                                                                                                                                                                    Claim 11; Page 174-175; 183pp; Japanese.
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HQYSKLPWT
                                     Similarity
9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 143-144; 183pp; Japanese
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                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             complementarity determining region;
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                                                 100.0%; Score 56; DB 4; Length 108; 100.0%; Pred. No. 0.043;
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Pred. No.
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RESULT 6
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RESULT 7
ABUILO13
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                               ABU11013 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Drugs containing genetically-modified antibody against ganglioside GD3 its fragment, immunocompetent cell activators or/and antitumor agents combination, applicable in treating malignant tumor like melanoma.
                               WO200278739-A1
                                                                                                                                                                                            Modified ganglioside GD3 antibody associated protein #6
                                                                                                                                                                                                                                                    04-FEB-2003
                                                                                                                                                                                                                                                                                                           ABU11013;
                                                                                 Mus musculus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABU11011 standard; protein; 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (KYOW ) KYOWA HAKKO KOGYO KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           89 HQYSKLPWT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HOYSKLPWT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HQYSKLPWT
                                                                                                                                        GD3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Niwa R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GD3;
                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                        anti-ganglioside
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    anti-ganglioside GD3 antibody; tumour; melanoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ø
                                                                                                                                                                                                                                                                                                                                                               protein;
                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kanazawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 56; DB 6;
Pred. No. 0.043;
Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ģ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ያ
                                                                                                                                           GD3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Asada
                                                                                                                                        antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                           tumour; melanoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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RESULT 8
AAR33257
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Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Drugs concernits fragment,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention describes drugs contain a genetically-modified antibody against ganglioside GD3 or its fragment and at least 1 of a substance capable of activating immunocompetent cells and a substance having an antitumor activity in combination. The drugs can be used to treat tumous like melanoma and can provide a treatment with enhanced therapeutic effect and little side-reaction, particularly to relieve problems of side-effects during the conventional single administration. This sequence represents a protein associated with the anti- ganglioside GD3 antibody
WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAR-2002;
                   Shitara K,
                                                                                                                           EP533199-A2
                                                                                                                                                         Protein
                                                                                                                                                                                         Key
                                                                                                                                                                                                              Rattus rattus
                                                                                                                                                                                                                                                                                    25-MAR-2003
12-JUL-1993
                                                                                                                                                                                                                                                                                                                     AAR33257;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAR-2001; 2001JP-00097483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-OCT-2002
                                                             18-SEP-1991;
                                                                                  18-SEP-1992;
                                                                                                       24-MAR-1993
                                                                                                                                                                              Peptide
                                                                                                                                                                                                                                  humanised;
                                                                                                                                                                                                                                                               Rat immunoglobulin L chain varible region of pKM641LA2
                                                                                                                                                                                                                                                                                                                                       AAR33257 standard; protein; 128 AA.
                                        (KYOW ) KYOWA
1993-095510/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2003-067410/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                containing genetically-modified antibody against ganglioside GD: ragment, immunocompetent cell activators or/and antitumor agents nation, applicable in treating malignant tumor like melanoma.
                                                                                                                                                                                                                                                                                                                                                                                            68
                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KYOWA HAKKO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ~
                                                                                                                                                                                                                                                                                                                                                                                                            HQYSKLPWT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 113; 121pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                            HQYSKLPWT 97
                                                                                                                                                                                                                                             variable; region;
                                                                                                                                                                                                                                 chimeric; antibody;
                   Hanai N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Niwa R,
                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                    (revised)
(first en
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                                       HAKKO KOGYO CO LTD
                                                             91JP-00238375
                                                                                  92EP-00116026
                                                                                                                                                        /note= "Signal peptide" 21. .128
                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                    entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kanazawa J,
                   Hasegawa M,
                                                                                                                                               "Mature
                                                                                                                                                                                                                                                                                                                                                                                                                                             .0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                           rat; immunoglobulin; heavy; H; chain;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Score
Pred.
                                                                                                                                              protein"
                                                                                                                                                                                                                                  expression vector
                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                   Miyaji H,
                                                                                                                                                                                                                                                                                                                                                                                                                                              No.
                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 6;
0.043;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                    Kuwana
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 108;
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RESULT 9
AAR53AS
AAR5ASAS
AC AAR5
AC A
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Best Local
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chimeric human antibody H chain expression. mRNA from mouse anti-GD3 monoclonal Ab KM641-producing cells was isolated and KM641 H and L chain cDNAs isolated. The base sequences of the Ig variable regions in KM641 H chain cDNA (pKM641HA3) and KM641 L chain cDNA (pKM641LA2) are given in AAQ45438-39. A KM641-derived chimeric human Ab H chain expression vector was constructed by joining the H chain variable region gene from pKM641HA3 to a vector for chimeric human Ab H chain expression using the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monoclonal antibody; Ab; ganglioside GM2; chimera; chimeric antibody; expression vector; heavy; light; chain; hypervariable region; CDR; constant region; hybridoma; Ig; immunoglobulin; promoter; enhancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 128 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The sequences given in AAR33256-57 represent rat heavy and light chain variable regions respectively. The DNA sequences encoding these proteins were used in the construction of humanised chimeric antibody expression vectors. In these humanised antibodies none of the amino acids of the non-human animal Ab variable region have been changed. (Updated on 25-MAR-new control of the co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Humanised chimeric antibody prodn. against ganglioside cancers, such as melanoma, neuroblastoma, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB; AAQ33258
                                                                                                                                                                                                                                                     Example 2 describes the construction of the vector pChi641HA1 for
                                                                                                                                                                                                                                                                                                                    Example 2; Page 115-116; 191pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Humanised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nakamura K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-MAR-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KM641 H chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR53340;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR53340 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (KYOW ) KYOWA HAKKO KOGYO KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1994-126857/16.
)B; AAQ45438.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   correct
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                                                                                                                                                                                                                                                                                                                                                                                                  antibody specific for ganglioside effect on cancers such as melanoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Koike M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92JP-00238452.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shitara
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ζ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hanai
                                                                                                                                                                                                                                                                                                                                                                                                  melanoma, neuroblastoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.051;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kuwana
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J
                                                                                                                                                                                                                                                                                                                                                                                                                                   for producing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hasegawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ı
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RESULT 11
AAB01628
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AC AAB01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 9
                                                                                                                  Matches
                                                                                                                            Query Match
Best Local (
                                                                                                                                                                                                                      This immunoglobulin region was isolated from pKM641LA2.A methionine codon, presumably the initiation codon ATG, was found in the vicinity of the 5' terminus and the sequence has a full length leader sequence. The chimeric human antibodies are useful in the treatment of cancer, especially that which is of neural ectodermal origin. In contrast to
                                                                                                                                                                              prior art constructs based on mouse monoclonal antibodies, the chimeric human antibodies do not cause anti-mouse immunoglobulin production. The chimeric human antibodies have a prolonged half-ilfe and a reduced frequency of adverse effects when compared to mouse monoclonal antibodies
  AAB01628;
                      AAB01628
                                                                                                                                                          Sequence 128 AA;
                                                                                                                                                                                                                                                                                                           Chimeric human antibody expression vectors.
                                                                                                                                                                                                                                                                                                                                          WPI; 1999-468416/39.
                                                                                                                                                                                                                                                                                                                                                                                                         07-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US5939532-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antibody; nucleotide; genomic; hypervariable region; chimeric;
light chain; amino acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pKM641 LA2 immunoglobulin light chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY28368;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY28368 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      synthetic DNAs given
                                                                                                                                                                                                                                                                                                                                                                                    (КҮОМ ) КҮОМА НАККО КОСУО КК
                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-AUG-1999
                                                                         109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 109
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                                                                                                                 Similarity 9; Conserv
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                                                                                         HQYSKLPWT
                                                                                                                                                                                                                                                                                                                                                               <u>,</u>,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HOYSKLPWT
                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HQYSKLPWT 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 128 AA;
                                                                         HOYSKLPWT 117
                                                                                                                                                                                                                                                                                     Col 99; 188pp; English.
                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                               Hanai N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry
                                                                                                                                                                                                                                                                                                                                                                                                         93US-00116778
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein; 128
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                    protein; 128
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                                                                                                                 Score 56; DB
Pred. No. 0.0
0; Mismatches
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                              Hasegawa M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 2;
0.051;
                                                                                                                          DB 2;
0.051;
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                                                                                                                                                                                                                                                                                                                                                               Koike M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 128
                                                                                                                                     Length 128;
                                                                                                                  Indels
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RESULT 12
AAB81994
ID AAB81
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Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                            The present sequence is a murine immunoglobulin light chain variable region from plasmid KM-641. The coding sequence was used in the creation of an expression vector, along with the sequence for a human antibody, to produce humanised chimaeric antibodies, which can be used to treat cancer. Humanised chimaeric antibodies are more effective than mouse antibodies as they do not provoke a reaction in the human and side effects, such as the formation of anti-mouse immunoglobulin antibody and the rapid half-life of the immunoglobulins, do not occur
                                                                  Ganglioside
                                                                                        03-JUL-2001
                                                                                                                                                                                                                                                                                         Sequence 128 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                            New humanized chimera antibody KM-871 useful for treating cancer, comprises variable region of mouse monoclonal antibody, reactive with ganglioside and human antibody constant region.
                                cancer.
                                            Ganglioside;
                                                                                                                AAB81994;
                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 14; Page 28-29; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Murine immunoglobulin
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                                                                                                                                     AAB81994 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EP1013761-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse; immunoglobulin; L'chain; light chain; variable region; cancer; humanised antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB; AAA51004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KYOW ) KYOWA HAKKO KOGYO KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-SEP-1991;
18-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Key
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                                                                                                                                                                                               HOYSKLPWT
                                                                                                                                                                                                                    HOYSKLPWT
                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hanai N,
                                                                   GD3
                                           GD3; complementarity determining region; CDR; antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91JP-00238375.
92EP-00116026.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99EP-00124345
                                                                  specific antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21. .128
/label= mature_immunoglobulin_light_chain_V_region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            l. .20
/label= signal_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                               117
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                                                                                                                                     protein;
                                                                                                                                                                                                                                                    100.0%;
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                                                                                                                                      128
                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                       Score 56; DB 3; Pred. No. 0.051;
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                                                                                                                                                                                                                                             Mismatches
                                                                   related protein #3.
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Synthetic

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RESULT 13
AAB81995
ID AAB81
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XX Gangl
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Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention
New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-SEP-1999; 99JP-00278291.
06-APR-2000; 2000JP-00105088.
                                                                            N-PSDB; AAF86901.
                                                                                                                                                 Hanai N,
                                                                                                                                                                                                                                          06-APR-2000; 2000JP-00105088
                                                                                                                                                                                                                                                                   30-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                    05-APR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                WO200123432-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ganglioside;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB81995 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a monoclonal antibody which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 155-156; 183pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB; AAF86898.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-SEP-2000; 2000WO-JP006774
                                                                                                                                                                                                                                                                                                                  29-SEP-2000; 2000WO-JP006774.
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                                                                                                                                                                                                (KYOW)
                                                                                                  2001-266143/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001-266143/27.
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Similarity 100.0%;
9; Conservative
                                                                                                                                                                                              KYOWA HAKKO
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                                                                                                                                            Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GD3
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                                                                                                                                                                                                                                                                      99JP-00278291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              specific antibody related protein #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               complementarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein; 128
                                                                                                                                                                                              KOGYO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KOGYO KK.
                                                                                                                                                 Nakamura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nakamura
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Pred. No. 0.0
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antibody;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
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RESULT 14
AAB81997
ID AAB81
XX AAB81
XX AAB81
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XX
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Query Match
Best Local S
Matches 9
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                                                                                                                                                                                                                   Sequence 128 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent
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06-APR-2000; 2000JP-00105088.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 161-162; 183pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     activity in cancer.
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DB; AAF86907.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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           Conservative
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                                                                                                         Length 128;
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RESULT 16
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            Synthetic
                                         Ganglioside;
                                                             Ganglioside
                                                                                   03-JUL-2001
                                                                                                       AAB81996;
                                                                                                                           AAB81996 standard;
                                                                                                                                                                                                                                                                Sequence 128 AA;
                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 140; 183pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                       activity in cancer.
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|HQYSKLPWT 117
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                                                             GD3
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                                        GD3; complementarity determining
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                                                             specific antibody related protein
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                                                                                                                          protein;
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RESULT 17
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06-APR-2000; 2000JP-00105088.
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                                                                        WPI; 2001-266143/27.
N-PSDB; AAF86913.
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06-APR-2000; 2000JP-00105088.
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                                                                                                                                                   Hanai N,
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ilarity 100.0%;
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Pred. No. 0.051;
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New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy

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06-APR-2000; 2000JP-00105088.
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                                                                                                              Sequence 128 AA;
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DB; AAF86895.
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06-APR-2000;
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Query Match
Best Local Similarity
Matches 9; Conserv
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DB; AAF86892.
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                                      GD3 specific antibody related protein #7.
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2000JP-00105088.
  GD3;
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complementarity determining region; CDR; antibody;
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Pred. No. 0.051;
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06-APR-2000; 2000JP-00105088
                     Drugs containing genetically-modified antibody against ganglioside its fragment, immunocompetent cell activators or/and antitumor ager combination, applicable in treating malignant tumor like melanoma.
                                                                                                                                                                                                                                                                                      29-MAR-2001; 2001JP-00097483
                                                                                                                                                                                                                                                                                                                                         29-MAR-2002; 2002WO-JP003170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU11003;
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                                                                                                                                                                                                                                   (KYOW ) KYOWA HAKKO
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                                                                                                                                                                          Niwa R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 anti-ganglioside GD3 antibody; tumour; melanoma.
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                                                                                                                                                                                                                                     KOGYO
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                                                                                                                                                                                  Kanazawa
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Pred. No.
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Best Local S
Matches 8
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Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention describes drugs contain a genetically-modified antibody against ganglioside GD3 or its fragment and at least 1 of a substance capable of activating immunocompetent cells and a substance having an antitumor activity in combination. The drugs can be used to treat tumour like melanoma and can provide a treatment with enhanced therapeutic effect and little side-reactions, particularly to relieve problems of side-effects during the conventional single administration. This sequence represents a protein associated with the anti- ganglioside GD3 antibody
                                                                                                                                                         The mouse VL gene product may be used to produce chimeric mouse- habs against HIV-1 comprising human Ig constant regions and murine variable regions. These novel sequence are useful in treatment, diand prophylaxis of HIV infections, and may be produced by a bacter yeast or mammalian expression system. (Updated on 25-MAR-2003 to correct PI field.)
                                                                                                              Sequence 126 AA;
                                                                                                                                                                                                                                                                                                                                                                                             New chimeric mouse human antibodies - used in treatment, diagnosis prophylaxis of HIV infections.
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19-AUG-1991
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                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 13; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; AAQ12017.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Better MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (XOMA ) XOMA CORP.
(GREC ) GREEN CROSS
(ZOMA-) ZOMA CORP.
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          Conservative
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100.0%; Pr
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                                Score 48;
Pred. No.
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Pred. No.
        Mismatches
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                                  DB 2;
1.3;
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                                                        Length 126;
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RESULT 24
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                                                                                                                                                                    Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                    This is the light (kappa) - chain variable (V) region of a mouse monoclonal antibody (MAb), 1C11, and is specific for an HIV-1 viral antigen. It is used in the construction of a chimeric MAb comprising heavy and light chains having murine V regions and human C regions. The chimeric MAbs are more effective than murine MAb 1C11 since they have an increased compatibility in humans. The heavy and light chain V-regions are joined by manipulating their respective joining (J) regions, to generate restriction enzyme recognition sites. The chimeric MAbs can be used as immunoconjugates, in association with e.g. toxins for HIV treatment. They can also be used in diagnosis of HIV. See also AAQ12056-60 and AAQ12052-G3. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1991-178105/24.
N-PSDB; AAQ12061.
          20-NOV-2003
                                                                                                                                                                                                                    Sequence 127 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                          New chimeric mouse-human antibodies --
-1 antigen from sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-NOV-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-MAY-1991.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chimeric antibodies; immunoconjugates; HIV; AIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Light (kappa) chain variable region of murine 1C11 immunoglobulin
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15-AUG-1991
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                                  ADA89188;
                                                         ADA89188 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Fig 13; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Better MD,
                                                                                                                                                                                                                                            correct DR field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (XOMA ) XOMA CORP.
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| QYSKLPWT 117
                                                                                                                      OYSKLPWT 117
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        (first entry)
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                                                        protein; 109 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein; 127
                                                                                                                                                                85.7%; but
100.0%; Pr
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                                                                                                                                                                               Score 48; DB 2; Pred. No. 1.3;
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                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                   CC domain. The protein binds a complex comprising a major histocompatibility complex (MHC) and a peptide, does not substantially bind the MHC in the complex (MHC) and a peptide, does not substantially bind the peptide in the absence of the MHC. The peptide is a peptide fragment of gpl10, CC (mUC1, TAX or hTERT. Also described: (1) a pharmaceutical composition (CC comprising the novel protein and a carrier; (2) a cytotoxic T cell (CC comprising one or more nucleic acids for expressing the Ig that binds a CC complex having an MHC and a peptide, does not substantially bind the MHC (CC in the absence of the bound peptide, and does not substantially bind the CC comprising a first segment that encodes the Ig variable domain; (4) a comprising a first segment that encodes the Ig variable domain; (4) a convel protein; (5) a transgenic animal whose genome includes heterologous nucleic acid sequences that encodes the protein; (6) identifying the CC novel protein; (5) a transgenic animal whose genome includes heterologous nucleic acid sequences that encodes the protein; (6) identifying the CC an antigen-binding protein; (8) ablating or killing a target cell that CC displays a peptide on a surface MHC molecule; (9) treating or preventing a cancerous disorder in a subject; and (10) detecting an MHC-peptide complex in a sample. A protein of the invention has cytostatic activity, CC and can be used in gene therapy. The protein is useful for preparing a CC complex in a sample. A protein of the invention has cytostatic activity, CC and complex the light chain of an antibody which binds to an MHC-peptide complex where the peptide component in as peptide fragment of epitide complex where the peptide component in as peptide fragment of component in 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New protein comprising an immunoglobulin heavy chain variable and an immunoglobulin light chain variable (VL) domain, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (DYAX-)
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light chain variable domain; major histocompatibility complex; MHC;
gp100; MUC1; TAX; hTERT; cytostatic; gene therapy; cancerous disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB; ADA89187.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hoogenboom HRJM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human antibody 2B2 light chain amino acid
                                                                                                                                                                            Sequence 109 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 7A; 224pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   preparing a composition for treating or preventing a cancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-FEB-2002; 2002US-0358994P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a protein comprising an immunoglobulin (Ig) heavy chain variable (VH) domain and an Ig light chain variable (VL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo
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                                           1 HOYSKLPWT
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  HQYGFLPWT 98
                                                                                       Conservative
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                                             9
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                                                                                                              Score 44;
Pred. No.
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RESULT 25 ABP43158 ID ABP43158 standard; protein; 215

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22-AUG-2002

(first entry)

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CC shock syndrome), inflammatory conditions (e.g., compenital and acquired (constitute), immune disorders (e.g., congenital and acquired immunedeficiencies, autoimmune ophoritis, systemic lupus erythematosus), cc immunodeficiencies, autoimmune ophoritis, systemic lupus erythematosus), cc immunodeficiencies, autoimmune ophoritis, systemic lupus erythematosus), cc respiratory disorders (e.g., anaemia), cardiovascular disorders, creative of the constant of the compounds which cc and urinary system disorders. Ovarian antigen polypeptides and cc polynucleotides may also be used in screening for compounds which cc modulate ovarian antigen expression or activity. The polynucleotides may further be used for gene therapy, chromosome mapping, in the cc identification of individuals and in forensic analysis, and the cc polypeptides may be used as food additives or to prepare antibodies conjectual in disease diagnosis, drug targeting and phenotyping. The present sequence represents a human ovarian antigen of the invention. Note: The sequence data for this patent did not form part of the printed conspectification, but was obtained in electronic format directly from WIPO as formulations.
                                                                                                                                                                                                                                                                                                                                                                                                                                          recombinant vectors and host cells comprising human ovarian antigen polynucleotides, antibodies against human ovarian antigens, and the use of ovarian antigen polynucleotides and polypeptides in diagnosing, treating, prognosing or preventing various ovary and/or breast-related disorders. Such conditions include ovarian cancer and breast cancer, and metastatic tumours of ovarian or breast origin, reproductive system disorders (e.g., infertility, disorders of pregnancy, anovulation, polycystic ovary syndrome, ovarian cysts, and dysmenorrhoea), endocrine disorders, infections (e.g., chlamydia, HI, toxoplasmosis, and toxic and dysmenorians).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ovarian antigen; ovary; ovarian; breast; cancer; tumour; ovarian cancer; breast cancer; tumour; reproductive system disorder; infertiality; pregnancy disorder; anovulation; polycystic ovary syndro; PCOS; ovarian cyst; dysmenorrhoea; endocrine disorder; infection; inflammatory condition; immune disorder; blood disorder; cardiovascular disorder; respiratory disorder; neurological disorder; gastrointestinal disorder; urinary system disorder; drug screening; gene therapy; chromosome mapping; forensic analysis; antibody preparation; cytostatic; immunomodulatory; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABP43228) and to cDNAs encoding them (ABQ54131-ABQ56305), and also encompasses polypeptides 90% identical and polynucleotides 95% identical to the sequences of the invention. The invention additionally relates to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated nucleic acid molecules encoding novel ovarian polypeptides, useful in the prevention, treatment and diagnosis of cancer (e.g. ovarian cancer), immune disorders, cardiovascular disorders and neurological
Sequence 215 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 11; SEQ ID NO 4290; 2922pp; English
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                                            ttp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention relates to 2175 novel human ovarian antigens (ABP41054-
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Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antibody chains, one or which supplies the catalytic protein, the invention provides for the binding specificity for a target protein, the invention provides for the construction of a repertoire of protein, the invention provides for the construction of a repertoire of protein with customised protein substrate specificities of potentially unlimited number and thus makes possible the effective treatment and/or prevention of any medical condition attributable to the presence or overexpression of an identified protein. The invention also provides mucleic acids encoding the catalytic antibodies (which can be used for gene therapy), host cells, transgenic non-human animals, and methods of cleaving a target protein (in vitro or in vivo) using a recombinant catalytic polypeptide. It also provides a library of recombinant catalytic polypeptides with altered enzymatic activity and a method of altering the enzymatic activity of the recombinant catalytic polypeptides with altered enzymatic activity and a method of altering the enzymatic activity of the recombinant catalytic polypeptides with altered enzymatic activity and a method of altering the enzymatic activity of the recombinant catalytic polypeptides.
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Key
Region
                                                                                                                                      Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel recombinant catalytic polypeptide useful for cleaving target proteins or for treating or preventing cancers, comprises a human antibody light chain operably joined to a heterologous antibody heavy
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                                                                                                                                      proteolytic A10-JK1 light chain, used in catalytic antibody
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The present sequence is the protein sequence of a human light chain variable region (VL), denoted A10-JK1, that is preferred for use in recombinant catalytic polypeptides the invention. VL genes encoding potentially catalytic variable regions were cloned by PCR from human genomic DNA following analysis of the human kappa repertoire for genes containing putative serine protease triads. Several genes were
                                                                                                                                                        Novel recombinant catalytic polypeptide useful for cleaving target proteins or for treating or preventing cancers, comprises a human antibody light chain operably joined to a heterologous antibody he
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containing putative serine identified, including Al0.
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                                                                                                                                                                Claim 2; Page; 33pp; English.
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The invention relates to a murine monoclonal antibody, MNAC13, which recognises the human nerve growth factor (NGF) receptor tyrosine kinase, TrkA, and acts as an antagonist of NGF/TrkA binding. MNAC13 has a light

The invention relates to a murine monoclonal antibody, MNAC13, which recognises the human nerve growth factor (NGF) receptor tyrosine kind TrkA, and acts as an antagonist of NGF/TrkA binding. MNAC13 has a light

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RESULT 29
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66.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ۲
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cc variable region (VL) as given in AAB48936, and a heavy chain cynthetic derivatives of monoclonal antibody MNAC13, such as the single chain antibody, scFvMNAC13 (AAB48937), which comprises the VL and VH cregions of MNAC13 in one polypeptide chain. The invention additionally contained to the invention comprises the VL and VH creates to pharmaceutical compositions comprising the antibody or cantibody derivatives of the invention, engineered eukaryotic cells able composition comprising the antibody or its derivatives, and a diagnostic composition comprising the antibody or its derivatives for in vivo compassition comprising the antibody or its derivatives for in vivo compassition comprising the antibody or its derivatives for in vivo compassition comprising the antibody and its derivatives for treating convological pathologies such as acute or chronic pain, neuromas, and convological pathologies such as acute or chronic pain, neuromas, and convological pathologies such as acute or chronic pain, neuromas, and convological pathologies such as acute or chronic pain, neuromas, and convological pathologies. The present sequence represents the anti-Trka single chain antibody scFvMNAC13
   ARBOULT 30
AABOG 354
ID AABOG
XX AABOG
XX AABOG
XX 17-OC
XX 17-OC
XX Cytoss
XW Cytoss
XW Cytoss
XW Cytoss
XW Compl
XX Co
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Best Local
The present sequence is the randomised VL3 region of a variant of antivascular endothelial growth factor (VEGF) antibody from phage library YV83. A large number of antibody variants containing randomised peptide inserts within the complementarity determining regions (CDRs) were
                                                                                                                                                                                                                                                                                      Antibody variants with higher binding affinity than native antibodies useful for diagnosis, prevention and treatment of neoplastic and non-neoplastic diseases comprises amino acid insertion in hypervariable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Humanised; antibody variant; phage display; randomised library; cytostatic; antiarthritic; antipsoriatic; antidiabetic; antiinflammatory; antiarteriosclerotic; vascular endothelia growth factor; VEGF; breast cancer; lung cancer; retinoblastoma; rheumatoid arthritis; psoriasis; atherosclerosis; diabetic retinopathy;
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                                                                                                                                                                                  Example
                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-387797/33
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                                                                                                                                                                                  Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence of anti-VEGF antibody variant Y0241-5.
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                                                                                                                                                                          58; 110pp; English
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66.7%;
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RESULT 31
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XX Human
KW Cytoss
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XX Homo
OS Synth
XX Homo
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Best Local S
Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                   Antibody variants with higher binding affinity than native antibodies useful for diagnosis, prevention and treatment of neoplastic and non-neoplastic diseases comprises amino acid insertion in hypervariable
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                                                                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Randomised VL3 sequence of anti-VEGF antibody variant Y0241-3.
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6; Conserv
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                                                                                                                                                                                                                                                                                                                Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0108945P
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                                                                                                                                                                                                                                                                                                                   58; 110pp;
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75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Muller
                                                                                                                                                                                                                                                                                                                   English.
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Pred.
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The present sequence is the randomised VL3 region of a variant of antivascular endothelial growth factor (VEGF) antibody from phage library YC 83. A large number of antibody variants containing randomised peptide inserts within the complementarity determining regions (CDRs) were prepared by phage display. Libraries were subjected to eight rounds of selection to isolate variants with an antigen binding affinity at least two-fold stronger than the binding affinity of parent antibody for the target VEGF antibody. The anti-VEGF antibody variants may be useful in diagnostic assays for detecting expression of VEGF in cells, tissue or serum. They may also be used in the prevention and treatment of

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RESULT 33
AAB62087
ID AAB62
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AC AAB62
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AAR30770
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Matches 6
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                                                                                                                                                                                                           The sequence is that of a consensus sequence hukl of the most abundant human subgroups, namely VL K 1 upon which 1s based the humanised variants of the light chain variable domain of murine anti-CD3 monoclonal antibody UCHT1 (muxCD3, Shalaby 1992). (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                             Sequence 107 AA;
                                                                                                                                                                                                                                                                                                                          WPI; 1993-018139/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Humanisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Consensus humanised murine anti-CD3 MAb UCHT1 VL domain hukI.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    neoplastic diseases such as breast cancer, lung cancer and retinoblastoma, and non-neoplastic diseases including rheumatoid arthritis, psoriasis, atherosclerosis, and diabetic and other proliferative retinopathies
            AAB62087;
                                                                                                                                                                                                                                                                  Disclosure; Fig 5; 126pp; English.
                                                                                                                                                                                                                                                                                       Humanisation of antibodies - by molecular modelling of the domains and alteration by gene conversion mutagenesis.
                                                                                                                                                                                                                                                                                                                                              Carter PJ,
                                                                                                                                                                                                                                                                                                                                                                                         14-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                               15-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                     23-DEC-1992.
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12-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR30770;
                                AAB62087 standard; protein; 107
                                                                                                                                                                                                                                                                                                                                                                    (GETH ) GENENTECH INC.
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                                                                                       90
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                                                                                                                                 6; Conserv
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                                                                                       QYNSLPWT
                                                                                                            QYSKLPWT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            QYSNVPWT
                                                                                                                                                                                                                                                                                                                                              Presta LG;
                                                                                                                                71.4%;
ilarity 75.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rapid; monoclonal antibody; muxCD3; light chain
                                                                                                                                                                                                                                                                                                                                                                                         91US-00715272
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                                                                                        97
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75.0%;
                                                                                                                                Score 40; DB:
Pred. No. 28;
1; Mismatches
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Pred. No. 1.7e+06;
1; Mismatches 1
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                                                                                                                                 1;
                                                                                                                                                     Length 107;
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                                                                                                                                                                                                                                                                                                    variable
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AAB60400
ID AAB6
XX
AC AAB6
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DT 24-A
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COnf
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COnf
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COnf
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COnf
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Anti
KW 1191
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                                                                                                                                                                                                            RESULT 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to treating cancer in a human susceptible to or chiagnosed with a disorder characterized by overexpression of Exb82 receptor. The method comprises administering an initial dose of at least approx.5 mg/kg of the anti-Erb82 antibody, and administering subsequent doses of the antibody in approximately the same amount or less than the initial dose. The method is useful for treating a disorder characterized by overexpression of Erb82 receptor, particularly a benign or malignant tumour, or a cancer. The cancer is selected from breast cancer tumour, or a cancer. The cancer is selected from breast cancer cancer, gastrointestinal cancer, non-small cell lung cancer, pancreatic cancer, gastrointestinal cancer, on-small cell lung cancer, ovarian cancer, iver cancer, bladder cancer, hepatoma, colon cancer, colorectal cancer, endometrial carcinoma, salivary gland carcinoma, kidney cancer, cancer, prostate cancer, vulval cancer, thyroid cancer, hepatic carcinoma and various types of head and neck cancer. The present sequence represents the human variable light chain consensus framework humkappal
                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 6
Anti-ErbB2 monoclonal antibody 2C4; HER2; humanised; VL; light chain variable region; cancer; cytostatic; EGFR-expressing cancer; epidermal growth factor receptor; colon cancer; rectal cancer; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Treating a human patient with a disorder characterized by overexpression of ErbB2 receptor such as a tumor or cancer (e.g. metastatic breast carcinoma) comprises administering several dosages of an anti-ErbB2
                                                                  Consensus human light chain kappa subgroup I framework, SEQ ID NO:5.
                                                                                                        24-APR-2001
                                                                                                                                         AAB60400
                                                                                                                                                                       AAB60400 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 107 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Fig 5A; 71pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-235058/24.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-AUG-1999; 99US-0151018P
23-JUN-2000; 2000US-0213822P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ErbB2 receptor; anti-ErbB2 antibody; tumour; cancer; breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                      (light kappa subgroup I)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            leukemia;
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                                                                                                                                                                                                                                                              OYNSLPWT
                                                                                                                                                                                                                                                                                               QYSKLPWT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cytostatic; carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                      (first
                                                                                                                                                                                                                                                                  97
                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                 71.4%;
75.0%;
                                                                                                                                                                                                                                                                                                                                 Score 40; DB
Pred. No. 28;
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                 Length 107;
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ARESULT 35
ARB61585
ID ARB61
XX
AC ARB61
XY
AC ARB61
XY
O1-AF
DT 04-AF
XX
XX
KW Erb82
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a method for treating cancer in a human patient, CC wherein the cancer expresses epidermal growth factor receptor (EGFR), AB60408). CC comprising administering an antibody which binds ErbB2 (HER2; AAB60408). CC In particular, the anti-ErbB2 antibody is the murine monoclonal antibody. CC AAB60396, AAB60397) or a humanised version of 2C4 (AAB60398. CC AAB60399). The invention also encompasses an isolated nucleic acid cencoding a humanised ErbB2-binding antibody; vectors and host cells comprising such nucleic acids; the recombinant production of a humanised CC emb2-binding antibody and a cytotoxic drug. The ErbB2-binding antibodies act by antagonising ErbB receptors, and as inhibitors of transforming CC growth factor alpha (TGF-alpha)-activated mitogen activated protein cespecially colon cancer, rectal cancer, colorectal cancer, lung cancer, cespecially colon cancer, rectal cancer, colorectal cancer (especially colon-small cell lung cancer), or breast cancer (especially colon-small cell lung cancer), or breast cancer (especially colorectal cancer). The antibodies may also have non-therapeutic colorectatic breast cancer is preferable to the use of EGFR-targetted CC uses e.g., as affinity purification agents. Using an antibody which binds to ErbB2 to treat cancer is preferable to the use of EGFR-targetted CC liver and skin, where the active drug will also bind, with skin toxicity for a present sequence represents the consensus human light chain kappa. CC subgroup I framework used to humanise the anti-ErbB2 antibody 2C4
                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 6
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                                     Human variable light chain consensus sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating cancer in a human, where the cancer expresses epidermal growth factor receptor (EGFR), comprises administering an antibody which binds
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ErbB2; cytostatic; prostate
                                                                                04-APR-2001
                                                                                                                       AAB61585;
                                                                                                                                                          AAB61585 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-080862/09.
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                                                                                                                                                                                                                                                             90 QYNSLPWT 97
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                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Fig 7A; 89pp; English
                                                                                                                                                                                                                                                                                                  QYSKLPWT 9
                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                (first entry
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75.0%;
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cancer; receptor tyrosine kinase; human;
                                                                                                                                                            107 AA
                                                                                                                                                                                                                                                                                                                                         Score 40; DB Pred. No. 28; 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                              4.
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RESULT 36
ADE71454
ID ADE71
XX ADE71
AC ADE71
XX 29-JA
DT 29-JA
XX Cytos
KW Cytos
KW ExbB2
KW neuro
KW adeno
KW squan
KW squan
KW squan
KW squan
KW squan
KW squan
KW caroc
KW cervi
KW cervi
KW cervi
KW breas
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KW breas
KW cervi
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The ErbB family of receptor tyrosine kinases are important mediators of cell growth, differentiation and survival. The receptor family includes four distinct members including Epidermal Growth Factor Receptor (EGFR or ErbB1), HER2 (ErbB2 or pl85 neu), HER3 (ErbB3) and Her4 (ErbB4 or tyro2). The present invention relates to a method for treating prostate cancer. The method comprises administering an antibody which binds ErbB2 and blocks ligand activation of an ErbB receptor. Preferably, the antibody blocks binding of monoclonal antibody 2C4 to ErbB2 and/or blocks TGF-alpha activation of mitogen-activated protein kinase (MAPK). The present sequence is the consensus framework for the human variable light chain. This sequence was used in a sequence alignment homology with the variable light chain of antibody 2C4 (see AAB61581)
                                                                               cytostatic; ErbB2 antagonist; tyrosine kinase inhibitor; cancer; ErbB2; ErbB2 antibody; carcinoma; lymphoma; blastoma; sarcoma; liposarcoma; neuroendocrine tumour; mesothelioma; schwanoma; meningioma; adenocarcinoma; melanoma; leukaemia; lymphoid malignancy; squamous cell cancer; epithelial squamous cancer; lung cancer; small-cell lung cancer; non-small cell lung cancer; lung cancer; adenocarcinoma of the lung; squamous carcinoma of the lung; squamous carcinoma of the lung; squamous carcinoma of the speritoneum; hepatocellular cancer; gastric cancer; glioblastoma; stomach cancer; gastrointestinal cancer; pancreatic cancer; bloader cancer; hepatoma; cervical cancer; oolon cancer; rectal cancer; colorectal cancer; breast cancer; colon cancer; rectal cancer; colorectal cancer; breast cancer; cancer; nectal cancer; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antibody; ErbB receptor; monoclonal antibody 2C4; variable light chain
                         thyroid
                            kidney cancer; renal cancer; prostate cancer; vulval cancer; thyroid cancer; hepatic carcinoma; anal carcinoma; penile ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human antibody variable light chain framework.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADE71454 standard; protein; 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 107 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Fig 7; 93pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Treating prostate cancer in a human comprises administering an antibody which binds ErbB2 and blocks ligand activation of an ErbB receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Agus DB,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         90 QYNSLPWT 97
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6; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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75.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
   anal carcinoma; penile carcinoma;
cer; tumour of the biliary tract;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4
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ADJ8808
ID BOOB
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XX CANCE
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OS Unide
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention describes a method of treating cancer that expresses ErbB2 comprising administering to a patient an antibody that binds ErbB2. Specifically claimed are antibodies that bind ErbB2, particularly monoclonal antibody 2C4 or humanised 2C4, or monoclonal antibody 4D5 or humanised 4D5. The methods are useful for treating cancer in a patient, particularly a human. The cancer includes carcinoma, lymphoma, blastoma, sarcoma, liposarcoma, neuroendocrine tumour, mesothelioma, schwanoma, meningioma, adenocarcinoma, melanoma, leukaemia, lymphoid malignancy, squamous cell cancer, non-small cell lung cancer, adenocarcinoma of the lung, squamous carcinoma of the lung, cancer, adenocarcinoma of the lung, squamous carcinoma of the lung, cancer of the peritoneum,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hepatocellular cancer, gastric or stomach cancer, gastrointestinal cancer, pancreatic cancer, glioblastoma, cervical cancer, ovarian cancer, liver cancer, bladder cancer, phatoma, breast cancer, colon cancer, rectal cancer, colorectal cancer, endometrial or uterine carcinoma, salivary gland carcinoma, kidney or renal cancer, prostate cancer, vulval cancer, thyroid cancer, hepatic carcinoma, anal carcinoma, penile cancer, testicular cancer, esophageal cancer, a tumour of the biliary tract, or head and neck cancer. This is the amino acid sequence of human antibody variable light chain framework that was used to produce a humanised antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Treating cancer (e.g. carcinoma, lymphoma or sarcoma) that expresses ErbB2 by administering to a patient an anti-ErbB2 antibody, and optionally an epidermal growth factor receptor-targeted drug or a tyrosine kinase inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-020226/02.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-JUN-1999; 99US-0141316P.
23-JUN-2000; 2000US-00602812.
 Homo sapiens
Unidentified
                                                         Cancer; ErbB2;
                                                                                           Human variable light chain consensus peptide
                                                                                                                               06-MAY-2004
                                                                                                                                                                    ADJ88008
                                                                                                                                                                                                    ADJ88008 standard; protein; 107 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 3; SEQ ID NO 5; 56pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US2003086924-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      head and neck cancer; human antibody; variable light chain framework.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                            2 QYSKLPWT 9
||: ||||
90 QYNSLPWT 97
                                                                                                                                                                                                                                                                                                90
                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                               Š
                                                                                                                             (first entry)
                                                       gene therapy; human
                                                                                                                                                                                                                                                                                                                                                                                        71.4%;
                                                                                                                                                                                                                                                                                                                                                                                      Score 40; DB
Pred. No. 28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English.
                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                         8
                                                                                                                                                                                                                                                                                                                                                                    1;
                                                                                                                                                                                                                                                                                                                                                                                                         Length 107
                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                    0;
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                                                                                                                       tunour, gastrinoma, islet cell cancer, mesothelioma, schwannoma, acoustic concuroma, meningioma, adenocarcinoma, melanoma, leukaemia, lymphoid comalignancy, squamous cell cancer, pithelial squamous cell cancer, lung cancer, small-cell lung cancer, non-small cell lung cancer, lung cancer, small-cell lung cancer, non-small cell lung cancer, lung cancer, small-cell lung, squamous carcinoma of the lung, cancer of the carcinoma, hepatocellular cancer, gastric or stomach cancer, gastrointestinal cancer, liver cancer, glioblastoma, cervical cancer, ovarian cancer, liver cancer, bladder cancer, hepatoma, breast cancer, colon cancer, rectal cancer, colorectal cancer, endometrial or cuterine carcinoma, salivary gland carcinoma, kidney or renal cancer, corcinoma, penile carcinoma, they cancer, hepatic carcinoma, anal carcinoma, penile carcinoma, testicular cancer, hepatic carcinoma, anal carcinoma, penile carcinoma, testicular cancer, osophageal cancer, a cumour of the biliary tract or head and neck cancer with anti-ErbB2 cantibodies. The invention is also useful in gene therapy. The present or sequence is human variable light chain consensus peptide.
                                           Matches
                                                           Query Match
Best Local S
                                                                                                                       Sequence 107 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-JUN-1999; 99US-0141316P.
23-JUN-2000; 2000US-00602812.
10-OCT-2002; 2002US-00268501.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        carcinoma, lymphoma, blastoma, medulloblastoma, retinoblast
liposarcoma, synovial cell sarcoma, neuroendocrine tumour,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3; SEQ ID NO 5; 56pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               to a patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Treating cancer that expresses ErbB2 e.g., breast, colon, rectal or colorectal cancer comprises administering an antibody that binds to ErbB2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-121529/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kelвey SM, Sliwkowski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JUN-2003; 2003US-00608626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US2004013667-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GETH ) GENENTECH INC.
                                       Similarity
6; Conserv
2 QYSKLPWT 9
                                         Conservative
                                                           71.4%;
75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               relates to methods for treating cancer such
blastoma, medulloblastoma, retinoblastoma, s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ₹
                                                           Score 40;
Pred. No.
                                             Mismatches
                                                               DB-8;
                                           ۲,
                                                                              Length 107;
                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                            toma, sarcoma, carcinoid
                                           0
                                           Gaps
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片
90 QYNSLPWT 97
                     0
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RESULT 38
ADN12054
ID ADN12
XX ADN12
XX I7-JU
DT 17-JU
DY Varia
XX tumor
XX tumor
XX Homo
OS Homo
XX W0200
XX W2020
XX W2020
XX I1-Jt
XX
                                                                                                                               Variable light chain consensus sequence
                                                             WO2004008099-A2
                                                                                  Homo sapiens
                                                                                                           tumor; anti-HER2 antibody; HER2/HER3; HER2/HER1; Cytostatic; cancer.
                                                                                                                                                           17-JUN-2004
                                                                                                                                                                                   ADN12054;
                                                                                                                                                                                                         ADN12054 standard; protein; 107
                                                                                                                                                           (first entry)
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11-JUL-2003; 2003WO-US021590

22-JAN-2004

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RESULT 39
ADP43328
ID ADP43
XX ADP43
XX ADP43
XX ADP43
XX DT 26-AU
XX Human
XX Human
XX Human
XX Erans
KW Epide
KW Chron
KW Chron
KW Gynae
KW Chron
XX Homo
OX Homo
XX Homo
XX
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treatment invention relates to identifying a tumor responsive to treatment with an anti-HER2 antibody involves detecting the presence of an HER2/HER3 and/or HER2/HER1 protein complex in a sample of the tumor. The method is useful in identifying a tumor as responsive to treatment with an anti-HER2 antibody. The methods and antibodies are useful in treating a condition or disorder including tumor or cancer, e.g. breast, prostate, lung, colorectal or ovarian cancer. The present accounts to the prostate of the present and the prostate of the present accounts.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chronic bronchitis; cystic fibrosis; cytostatic; antipsoriatic; gynaecological; dermatological; vasotyppic; antiarteriosclerotic; cardiant; antianginal; antilipaemic; hypotensive; antiasthmatic; antiallergic; antiinflammatory; antitussive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   epidermal growth factor receptor; ErbB; ErbB2;
transforming growth factor alpha; benign hyperroliferative disorder;
psoriasis; endometriosis; scleroderma; vascular disease; atherosclerosis;
restenosis; colon polyps; fibroadenoma; respiratory disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Koll
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JUL-2002;
20-JUN-2003;
  Treating non-malignant endometriosis, involvir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 5; 105pp; English.
                                                                                     WPI; 2004-450361/42.
                                                                                                                                           Sliwkowski
                                                                                                                                                                                                                                                    21-NOV-2002; 2002US-0428027P
                                                                                                                                                                                                                                                                                                          21-NOV-2003; 2003WO-US037367
                                                                                                                                                                                                                                                                                                                                                                                                                      WO2004048525-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human; variable light kappa chain; antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADP43328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADP43328 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  represents a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HER2/HER3 and/or HER2/HER1 protein complex.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying a tumor responsive to treatment with an anti-HER2 antibody
                                                                                                                                                                                                                                                                                                                                                                   10-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GETH )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2004-156546/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  monoclonal variable light kappa chain antibody SeqID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90
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                                                                                                                                                                                                 GENENTECH
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م
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QYNSLPWT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                QYSKLPWT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bossenmaier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treating cancer, comprises detecting the
                                                                                                                                           ₹
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002US-0396290P.
2003US-0480043P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  variable light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
llignant disease or disorder such involving abnormal activation or
                                                                                                                                           Brunetta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             71.4%;
75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B
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                                                                                                                                           g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chain consensus sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 40; DB
Pred. No. 28;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 107;
as psoriasis, production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 presence of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kelsey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>.</u>
     epidermal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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growth factor receptor or ErbB ligand by binds ErbB2 to mammal.
                                                                                           Example 3; SEQ ID NO 5; 74pp; English
                                                                                                         administering antibody that
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benign hyperproliferative disorder, psoriasis, endometriosis, scleroderma, vascular disease (such as atherosclerosis or restenosis), colon polyps, fibroadenoma or respiratory disease (such as chronic bronchitis or cystic fibrosis). Accordingly, they exhibit cytostatic, antipsoriatic, gynaecological, dermatological, vasotropic, antiarteriosclerotic, cardiant, antianginal, antilpsemic, hypotensive, antiasthmatic, antiallergic, antiinflammatory and antiatussive activities. This polypsptide sequence is the human variable light kappa chain protein used to humanise the murine anti-ErbB2 proteins of the invention. This invention refers to a novel method for treating a non-malignant disease that involves abnormal activation or production of an epidermal growth factor receptor (ErbB) receptor or an ErbB ligand. Specifically, it refers to the use of humanised murine anti-ErbB2 antibodies to block ligand activation of the ErbB receptor, where the ErbB ligand is a transforming growth factor alpha that promotes mitogen-activated protein kinase (MAPK). The present invention describes conjugating the monoclonal murine antibody 2C4, or more particularly the humanised antibody 574, to a cytotoxic or therapeutic agent such that it can be used to treat a cytotoxic or therapeutic agent such that it can be used to treat

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Best Loc
Matches
                                    Query Match
Best Local
90
              2 QYSKLPWT 9
                            Similarity 6; Conser
QYNSLPWT
                            Conservative
 97
                                    71.4%;
75.0%;
                            1:
                                    Score 40;
Pred. No.
                            Mismatches
                                    DB
28;
                                           8
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                                           Length 107;
                            0
                            Gaps
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Sequence 107 AA;

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ARESULT 40
AAW04177
ID AAW04
AC AAW04
A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9629350-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein; 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of
E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human; Fas ligand; monoclona
apoptosis; assay; diagnosis;
leosis; systemic lupus erythe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fas ligand antibody NOK-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ξ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     monoclonal; antibody;
diagnosis; disease;
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Claim 20;

Page 80-81; 133pp; Japanese

Kayagaki N,

Yagita

H,

Okumura

Nakata

1996-443140/44.

AAT39550

(SUME)

SUMITOMO ELECTRIC IND

8 ζ,

20-MAR-1995; 27-OCT-1995; 21-MAR-1996; 26-SEP-1996

95JP-00087420. 95JP-00303492. 96WO-JP000734

Monoclonal antibody specifically recognising the Fas ligand - useful for the detection of Fas ligands either on cell surface or in solution.

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CC The present sequence is a variant light chain variable region of the anticle chuman Fas ligand monoclonal antibody (MAD) NOK-1. NOK-1 is produced by the hybridoma NOK-1 (FERM BP-5044), which was prepared by immunising mice with transformed human Fas ligand expressing COS cells isolated from the mice with myeloma P3x63Ag8.653 (ATCC CRL-1580) cc cells. The MAD recognises the human Fas ligand on the cell surface or in continuous and can be used to inhibit the apoptosis inducing cell surface or in the surface or in continuous cells, and can be used for a Fas ligand assay consists, e.g. hepatitis, infectious mononucleosis and systemic lupus xx sequence 108 AA;

Query Match Sequence 118, Mismatches 1; Indels 0; Gaps 0;

Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Search completed: December 17, 2004, 18:29:07

Job time: 29.3539 secs
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Sequence 52477, A
Sequence 18, Appl
Sequence 18, Appl
Patent No. 5200183
Patent No. 5215909
Sequence 1, Appli
Sequence 2, Appli
Sequence 3, Appli
Sequence 5, Appli
Sequence 6, Appli
Sequence 6, Appli
Sequence 7, Appli
Sequence 9, Appli
Sequence 10, Appli
Sequence 11, Appli

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein search,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Issued Patents AA:*

1: /cgn2_6/ptodata/1

2: /cgn2_6/ptodata/1

3: /cgn2_6/ptodata/1

4: /cgn2_6/ptodata/1

5: /cgn2_6/ptodata/1

6: /cgn2_6/ptodata/1
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/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/backfiles1.pep:*
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US-09-225-322B-10
US-09-264-304-110
US-09-764-304-110
US-09-764-304-126
US-09-440-781-26
US-09-440-781-26
US-09-440-781-26
US-09-440-781-26
US-09-440-781-26
US-09-705-686-18
US-09-705-686-18
US-09-705-392A-18
US-09-705-392A-18
US-09-705-089-3
US-08-974-899-3
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US-09-440-781-25
US-09-440-781-25
US-09-440-781-25
US-09-440-781-94
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Compugen Ltd
Sequence 10, Appl
Sequence 19, Appl
Sequence 19, Appl
Sequence 24, Appl
Sequence 26, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 14, Appl
Sequence 14, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 3, Appli
Sequence 25, Appl
Sequence 27, Appl
Sequence 21, Appl
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Fatent NO. 6437098

GENERAL INFORMATION:

APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: KUWANA, YOSHHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101

CURRENT APPLICATION NUMBER: US/09/225,322B

CURRENT FILING DATE: 1995-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1991-09-18

NUMBER OF SEQ ID NO 10
SEQ ID NO 
  US-09-25-322B-19
US-09-25-322B-19
; Sequence 19, Application US/09225322B
; Patent No. 643709B
; Patent No. 643709R
; GENERAL INFORMATION:
; APPLICANT: SHITARA, KENYA
; APPLICANT: HANAI, NOBUO
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US-09-225-322B-10
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Best Local S
Matches 9
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109 HQYSKLPWT 117
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US-08-797-689-18
US-09-984-186-18
5205183-5
5215909-12
US-08-446-100-1
US-08-446-100-3
US-08-446-100-4
US-08-446-100-6
US-08-446-100-6
US-08-446-100-7
US-08-446-100-9
US-08-446-100-9
US-08-446-100-10
US-08-446-100-11
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US-08-446-100-13
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Pred. No.
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0.0042;
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Result No.

Score

Query Match

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Gaps

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Minimum Maximum

DB 80

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length:

Total number

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Sequence: Title: Perfect score: OM protein -

Scoring table:

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; OTHER INFORMATION: cDNA KM-641 US-09-764-304-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-764-304-10
                                                                                                                                                                                                                                CURRENT FILING DATE: 199-00-18
EARLIER FILING DATE: 2001-01-19
EARLIER FILING DATE: 2001-01-09-225,322
EARLIER FILING DATE: 1999-01-05
EARLIER FILING DATE: 1999-01-05
EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER FILING DATE: 1995-03-21
EARLIER FILING DATE: 1994-08-17
EARLIER FILING DATE: 1994-08-17
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER FILING DATE: 1992-09-17
EARLIER FILING DATE: 1992-09-17
EARLIER FILING DATE: 1992-09-18
EARLIER FILING DATE: 1991-09-18
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CURRENT APPLICATION NUMBER: US/09/225,322B
CURRENT FILING DATE: 1999-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1994-08-17
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SEQ ID NO 19
LENGTH: 128
                                                                                                                                                            SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
                                                                                                                                                                                        SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US07/947,674
PRIOR FILING DATE: 1992-09-17
PRIOR APPLICATION NUMBER: UP 3-238375
PRIOR FILING DATE: 1991-09-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 249-101
                                                     FEATURE:
                                                                      ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
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                                                                                                           TYPE: PRT
                                                                                                                                     ENGTH: 128
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9; Conserv
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MIYAJI, HIROMASA
KUWANA, YOSHIHISA
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APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: HYSAJI, HIROWASA
APPLICANT: KUYANA, YOSHIHISA
APPLICANT: KUWANA, YOSHIHISA
FILE REFERENCE: 249-101
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Best Local S
Matches 9
                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Yvonne Man-yee Chen et al.
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: p1469R1
                                                                                                      CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
SEQ ID NO 24
                                                                                                                                                                                                                                                 Sequence 24, Application US/09440781 Patent No. 6632926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 19
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Best Local (
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NUMBER OF SEQ ID NOS: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EARLIER APPLICATION NUMBER: US07/947,674
EARLIER FILING DATE: 1992-09-17
EARLIER APPLICATION NUMBER: JP 3-238375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER FILING DATE: 1994-08-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER FILING DATE: 1999-01-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER FILING DATE: 1995-05-31
                                                                     LENGTH: 9
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
NAME/KEY: artificial LOCATION: 1-9
                                   FEATURE:
                                                  ORGANISM: artificial sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 128
                                                                                                                                                                                                                                                                                                                                                            109 HQYSKLPWT 117
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o. 6495666
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; NAME/KEY: artificial
; LOCATION: 1-9
; OTHER INFORMATION: variant CDR sequence
US-09-440-781-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: variant CDR sequence US-09-440-781-24
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                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 7
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Best Local Similarity /3...
6; Conservative
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SEQ ID NO 26
LENGTH: 9
TYPE: PRT
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Best Local (
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Patent No. 663292
                                                                                                                                                                                                                                                                                                                                                                           Sequence 18, Appli
Patent No. 5821337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: P1469R1
CURRENT APPLICATION NUMBER: US/09/440,781
CURRENT FILING DATE: 1999-11-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Yvonne Man-yee Chen et al.
TITLE OF INVENTION: ANTIBODY VARIANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: artificial sequence
                                                                                                                                               SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                  APPLICANT: Leonard G. Presta
TITLE OF INVENTION: Immunoglobulin Variants
          FILING DATE: 15-JUN-
PRIOR APPLICATION DATA:
                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/05126
                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                   APPLICATION NUMBER: US/07/934,373C FILING DATE: 21-Aug-1992 CLASSIFICATION: 530
                                                                                                                                           OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                       ADDRESSEE:
 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                              INFORMATION:
                                                                                                                                                                                                    94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 OYSSVPWT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 QYSKLPWT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 6; Conserv
                                                                                                                                                                                                                                            South San Francisco
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                                                                                                                                                                                                                              California
                                                                                                                                                                                                                                                                                                                                                                                       Application US/07934373C
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                                                                                                                                                                                                                                                         1 DNA Way
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                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                Paul J.
                                                                                                                                                                                                                                                                     Genentech, Inc.
                            15-JUN-1992
                                                                                                                                                                                                                                                                                                                                             Carter
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75.0%;
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75.08;
07/715272
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Pred. No. 3.8e+05;
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US-07-934-373C-18
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                                               INFORMATION FOR SEQ ID NO: 18:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tent No. 6054297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 107 amino acids
TYPE: Amino Acid
                               SEQUENCE CHARACTERISTICS
                                                                        TELEPHONE: 650/225-1994
                                                                                                                                                    FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                               APPLICATION NUMBER: PCT/US92/05126
FILING DATE: 15-JUN-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 17-NOV-
                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US.
                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Immunoglobulin Variants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity nes 6; Conserv
TYPE: Amino Acid
                                                                                                          REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: PO
                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: WinPatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: UZIP: 94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: South San
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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1 DNA Way
                                                             650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Leonard G. Presta
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                                                                                                                                      Wendy M.
                                                                                                                                                                      UMBER: 07/715272
14-JUN-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                  (Genentech)
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Pred. No.
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Query Match
Best Local Similarity
""" hes 6; Conserv:
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US-08-146-206C-18
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                                                                                                                  Sequence 14, Application US/09648067A Patent No. 6627196 GENERAL INFORMATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WINDRALIN (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/146,206C
PILING DATE: 17-No. 6407213-1993
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/715272
PILING DATE: 14-UN-1991
APPLICATION NUMBER: 07/715272
PILING DATE: 14-UN-1991
ATTORNEY/AGENT INFORMATION:
APPLICANT: Baughman, Sharon A.
APPLICANT: Shak Steven
TITLE OF INVENTION: Dosages for Treatment with Anti-ErbB2 Antibodies
FILE REFERENCE: P1775R1
CURRENT APPLICATION NUMBER: US/09/648,067A
CURRENT FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/151,018
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CITY: South San Fra
CTATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: PO-
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Carter, APPLICANT: Presta,
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                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
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                                                                                                                                                                                                                                             90 QYNSLPWT 97
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Similarity 75.0%;
6; Conservation
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1 DNA Way
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Leonard G.
Method for Making Humanized Antibodies
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75.0%;
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US-09-705-686-18
                                                                                                           US-09-705-686-18
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Patent No. 6639055
GENERAL INFORMATION:
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Matches
                                    Matches
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LENGTH: 107
                                                    Query Match
Best Local (
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PRIOR APPLICATION NUMBER: US 60/213,822
PRIOR FILING DATE: 2000-6-23
NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial sequence
                                                                                                                                                                                          TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
APPLICATION NUMBER: 07/715272
FILING DATE: 14-UN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: 90709P1D3
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
TRILFBRN'. 650/225-1994
TRILFBRN'. 650/225-1994
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nes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/705,686
FILING DATE: 02-No. 6639055-2000
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AFFLICANT: Carter, Paul J.

Presta, Leonard G.
TITLE OF INVENTION: Method for Making Humanized Antibodies
NUMBER OF SEQUENCES: 26
                                                                                                                        TYPE: Amino Acid
TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 i
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ADDRESSEE: Genentech, Inc.
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   2 QYSKLPWT 9
                                    Similarity 6; Conserv
                                                                                                                                                                             LENGTH: 107 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: California
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                                        Conservative
                                                    71.4%;
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                                      Score 40; DB 4
Pred. No. 3.4;
1; Mismatches
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Pred. No.
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                                                                                                                              18:
                                                                       DB 4; Length 107;
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US-09-705-392A-18
                                                                                                                                                                                                                                                                                                    RESULT 13
PCT-US93-07832-18
                                                                                                                                                                                                                                                          Sequence 18, Application PC/TUS9307832 GENERAL INFORMATION:
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 18:
COMPUTER READABLE FORM:
MEDIUM TYPE: · 5.25 inch, 360 Kb floppy disk
COMPUTER: IBM PC compatible
                                                                                              CORRESPONDENCE ADDRESS: Inc.
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                                                                                                                                                                                TITLE OF INVENTION: Immunoglobulin Variants
                                                                                                                                                                                                                                            APPLICANT:
                                                              COUNTRY:
ZIP: 940
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 / Match 71.4%;
Local Similarity 75.0%;
les 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
APPLICATION NUMBER: 07/715272
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/705,392A
FILING DATE: 02-No. 6719971-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 18:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Presta, Leonard G.
TITLE OF INVENTION: Method for Making Humanized Antibodies
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                     90 QYNSLPWT 97
                                                                                                                                                                                                                                                                                                                                                                                                                           2 QYSKLPWT 9
                                                              94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 107 amino acids TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: P0709P1D1 REVISED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Lee,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TY: South San Francisco
                                                                                                                                                                                                                                       Genentech, Inc.
                                                                                 USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 40; DB 4; Length 107; Pred. No. 3.4; Indels 1; Indels
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/031971
PILING DATE: 11/27/96
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: P1014R1
TELECOMMUNICATION; INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Presta, Leonard G.
APPLICANT: Jardieu, Paula M.
TITLE OF INTENTION: Humanized Anti-CD11a Antibodies
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US9
FILING DATE: 19930820
                                                                                                                                                                                                            SOFTWARE: WinPatin (Ge
CURRENT APPLICATION DATA
                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/934373
FILING DATE: 21-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 07/7.
FILING DATE: 14-JUN-1991
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                 STREET: 1 DNA "",
CITY: South San Francisco
STATE: California
                                                                                                                                                      APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                 ZIP: 94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
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TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: PCT/
FILING DATE: 15-JUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                           DDRESSEE:
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75.0%;
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                                                                                                                                                                                                                          (Genentech)
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Pred. No.
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RESULT 16
US-09-795-798-3
; Sequence 3, Application US/09795798
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Best Local Similarity 75.0
Whiches 6; Conservative
                                                                                                                                        Matches
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                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                           FILING DATE:

FILING DATE:

CLASSIFICATION: 424

ATTORNEY/AGENT INFORMATION:

NAME: BUCCA Ph.D., Daniel

REGISTRATION NUMBER: P-42,368

REGISTRATION NUMBER: 5035

TELECOMMUNICATION INFORMATION:

TELEPHONE: 703-518-5100

TELEPAX: 703-684-1124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS: LENGTH: 108 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                            TOPOLOGY: li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: OKUMURA, KO
APPLICANT: NAKATA, MOEOMİ
TITLE OF INVENTION: THERAPEUTIC AGENT FOR HEPATITIS
NUMBER OF SEQUENCES: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                       Local Similarity hes 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Alexandria
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 22314
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                                                                                                                                                                                                                                                       i: 108 amino acids amino acid
                                                                              QYSEFPWT 97
                                                                                                          QYSKLPWT 9
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YAGITA, Hideo
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                                                                                                                                        Conservative
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75.0%;
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75.0%;
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Pred. No. 3.4;
1; Mismatches
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                                                                                                                                                                  DB 3;
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GENERAL INFORMATION:
APPLICANT: Presta, Leonard G.
Jardieu, Paula M.
Jardieu, Paula M.
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 31, Application US/08560558E Patent No. 5891996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local :
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: WINDOWS95
SOFTWARE: WORDPETECT 5.1/5.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/560,59
                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: ITILE OF INVENTION: ITILE OF INVENTION: INVENTION: INVENTION: IT
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                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/974,899
FILING DATE: CUNKNOWN
FILING DATE: CUNKNOWN
ATTORNEY,AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: P1014R1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
TELEPHONE: 650/225-1981
                                                                                                                                             COUNTRY: UI
                                                                                                                                                                                     STATE: Utah
                                                                                                                                                                                                       STREET: P.O. Box 225
CITY: Salt Lake City
                                                                                                                                                                                                                          ADDRESSEE: Allen C. Turner, TRASK, BRITT & ROSSA STREET: P.O. Box 2250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/795,798
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
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TITLE OF INVENTION: Humanized Anti-CD11a Antibodies
NUMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
                                                                                                                                                                  United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      71.4%;
75.0%;
                                                                                                                                                                                                                                                                                                        Humanized and chimeric monoclonal antibodies that recognize epidermal growth factor receptor EGF-R); diagnostic and therapeutic use.
                                                                                                                                                                                                                                                                                          34
    US/08/560,558E
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                                                                                                                                                                  of America
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; TYPE: PRT
; ORGANISM: Murine
; FEATURE:
; NAME/KEY: MISC FEATURE
; OTHER INFORMATION: CDR c
US-09-217-268B-31
                                                                                                                                      US-09-440-781-25
Sequence 25, Application US/09440781
Patent NO. 6632926
GENERAL INFORMATION:
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Matches
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Best Local :
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SEQ ID NO 25
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Humanized and Chimeric Monoclonal Antibodies That Recognize Epide TITLE OF INVENTION: Growth Factor Receptor (EGF-R); Diagnostic and Therapeutic Use FILE REFERENCE: 2720.1US
CURRENT APPLICATION NUMBER: US/09/217,268B
CURRENT FILING DATE: 1998-12-21
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn version 3.1
                                               APPLICANT: Yvonne Man-yee Chen et al. TITLE OF INVENTION: ANTIBODY VARIANTS FILE REFERENCE: P1469R1 CURRENT APPLICATION NUMBER: US/09/440,781 CURRENT FILING DATE: 1999-11-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Mateo de Acosta del Rio, Christina M
APPLICANT: Rodriquez, Rolando P
APPLICANT: Frias, Ernesto M
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: No. 5891996ember 17, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 9 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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REGISTRATION NUMBER:
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nilarity 75.0%;
Conservative
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ilarity 75.0%;
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Pred. No. 3.8e+05;
1; Mismatches 1
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Pred. No. 3.8e+05;
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; FEATURE;
; NAME/KEY: artificial
; LOCATION: 1-110
; OTHER INFORMATION: humanized antibody light chain variable domain
US-09-440-781-95
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                         Query Match
Best Local S
Matches 6
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CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
SEQ ID NO 94
                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: YVONNE Man-yee Chen et al.
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1
                                                                                                                                                                                                                                                                                                                  Sequence 95, Appli
Patent No. 6632926
                                                                                                                                                                                    SEQ ID NO 95
LENGTH: 110
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Matches 6; Conserv
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Patent No. 6632926
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Best Local
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CURRENT FILING DATE: 1999-11-16
NUMBER OF SEQ ID NOS: 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Yvonne Man-yee Chen et al. TITLE OF INVENTION: ANTIBODY VARIANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 110
TYPE: PRT
ORGANISM: artificial sequence
FEATURE:
NAME/KEY: artificial
LOCATION: 1-110
COTHER INFORMATION: humanized antibody light chain variable domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: P1469R1
                                                                                                                                                         ORGANISM: artificial
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ORGANISM: artificial sequence
FEATURE:
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6; Conserv
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2 QYSKLPWT 9
                           Similarity 6; Conserv
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75.0%;
                                        Score 39;
Pred No.
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Pred. No.
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Pred. No. 3.8e+05;
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                                                       Length 110
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US-08-560-558E-27
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US-08-497-312-18
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                                                                                                                                                                               Patent No.
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APPLICANT:
                                                APPLICANT:
TITLE OF INVENTION: Huma
TITLE OF INVENTION: anti
TITLE OF INVENTION: EGF-
NUMBER OF SEQUENCES: 34
CORRESPONDENCE ADDRESS:
                                                                                                                                                               GENERAL INFORMATION:
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TELEX: 388961 1PM04UT
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: prote
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PATENTIN Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/497,312
FILING DATE: 30-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: CU 80/94
FILING DATE: 30-JUN-1994
ATTORNEY/AGENT INFORMATION:
NAME: BOND, LAURENCE B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TITLE OF INVENTION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 801/532-1922
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ADDRESSEE: Allen C. Turner, TRASK, BRITT & ROSSA STREET: P.O. Box 2250 CITY: Salt Lake City
                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                no protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      unknown
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75.0%;
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                                                                                      Humanized and chimeric monoclonal antibodies that recognize epidermal growth factor receptor EGF-R); diagnostic and therapeutic use.
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Pred. No. 5
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5.6;
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; HYPOTHETICAL:
US-08-560-558E-27
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                                                                                                                           US-09-217-268B-27
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INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 114 amino
                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/217,268B
CURRENT FILING DATE: 1998-12-21
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin version 3.1
SEQ ID NO 27
                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 27, Application US/09217268B Patent No. 6506883
                                                                          Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                            Matches
                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Humanized and Chimeric Monoclonal Antibodies That Recognize Epider TITLE OF INVENTION: Growth Factor Receptor (EGF-R); Diagnostic and Therapeutic Use FILE REFERENCE: 2720.1US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: WINDOWS95
SOFTWARE: WORDERFIECT 5.1/5.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/560,5.
FILING DATE: No. 5891996ember 1
ATTORNEY/AGENT INFORMATION:
NAME: Turner, Allen C.
REGISTRATION NUMBER: 33,041
                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Mateo de Acosta del Rio, Christina M
APPLICANT: Rodriquez, Rolando P
APPLICANT: Frias, Ernesto M
                                                                                                                                                                    TYPE: PRT
ORGANISM: Murine R3 antibody
FEATURE:
                                                                                                                                        NAME/KEY: MISC_FEATURE OTHER INFORMATION: Ded
                                                                                                                                                                                                                     LENGTH:
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TELECOMMUNICATION INFORMATION:
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TOPOLOGY: unl
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95 QYSHVPWT 102
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                                                                                                                                          Deduced amino acid sequence of VK of murine
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75.0%;
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                                                                             Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 39; [
Pred. No. 5.
                                                              Mismatches
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                                                                                            DB 4;
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                                                                                        Length 114;
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                                                              Gaps
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                                                                                                                                           antibody
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RESULT 25 US-09-217-268B-35 ; Sequence 35, Application US/09217268B

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US-08-497-312-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Humanized and Chimeric Monoclonal Antibodies That Recognize Epide TITLE OF INVENTION: Growth Factor Receptor (EGF-R); Diagnostic and Therapeutic Use FILE REFERENCE: 2720.1US (CURRENT APPLICATION NUMBER: US/09/217,268B CURRENT FILING DATE: 1998-12-21 NUMBER OF SEQ ID NOS: 36
SOPTWARE: Patentin version 3.1
SEQ ID NO 35
LENGTH: 114
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Amino acid sequence of humanized VK of murine R3 antibody contain OTHER INFORMATION: ing mAb CDRs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 14, Application US/08497312
Patent No. 5712120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Mateo de Acosta del Rio, Christina M
APPLICANT: Rodriquez, Rolando P
APPLICANT: Frias, Ernesto M
                                                                                                                                                                                                                                   PRILING DATE: 30-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: CU 80/94
FILING DATE: 30-JUN-1994
ATTORNEY/AGENT INFORMATION:
NAME: BOND, LAURENCE B.
REGISTRATION NUMBER: 30,549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                      TELEX: 388961 1PM04UT
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
LENGTH: 116 amino aci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: 1
TITLE OF INVENTION: 1
TITLE OF INVENTION: 6
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/497,312
FILING DATE: 30-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 26
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
les 6; Conserv
                                                       TYPE: amino acid
                                                                                                                                                      TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
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                                                                                                                                                                            TELEPHONE:
                               STRANDEDNESS:
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                                                                        116 amino acids
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                 unknown
                                                                                                                                                                                801/532-1922
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                                    unknown
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75.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Method for obtaining modified immunoglobulins with reduced immunogenicity of murine antibody variable domains, compositions containing them.
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Pred. No. 5
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Query Match
Best Local Similarity
Tatches 5; Conserve
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US-09-270-767-52477
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US-09-270-767-37260
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APPLICANT: Homburger et al.

APPLICANT: Homburger et al.

TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster

TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster

TITLE OF INVENTION: NUMBER: US/09/270,767

CURRENT FILING DATE: 1999-03-17

NUMBER OF SEQ ID NOS: 62217

SOFTWARE: Patentin Ver. 2.0

LENGTH: 191

TYPE: PRT
                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Homburger et al.
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 52477
                                                                                                                                                                                                                                                                                                                                                   Sequence 52477, Application US/09270767 Patent No. 6703491
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Best Local
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Best Local
                                                                                                                                                   ORGANISM: Drosophila melanogaster FEATURE:
                                                                                                                                                                                   TYPE: PRT
                                                                                                                                     OTHER INFORMATION: Xaa means any amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Xaa means any amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Drosophila melanogaster
                                                                                                                                                                                                LENGTH: 191
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nes 5; Conserv
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HRYSQRPWS
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                                                            Conservative
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75.0%;
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55.6%;
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Pred. No.
                                                                          Score 38;
Pred. No.
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Pred. No.
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                                                                       DB
15;
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                                                                                       Length 191;
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RESULT 29 US-08-797-689-18

; Sequence 18, Application US/08797689

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RESULT 30
US-09-984-186-18
; Sequence 18, Application
; Patent No. 6686179
; GENERAL INFORMATION:
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Best Local Similarity
Matches 6; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-797-689-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: System 7.1
SOFTWARE: Word 5.1 (Patentin)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Fleer,
APPLICANT: Fourni
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/797,689
FILING DATE: 31-JAN-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/256,927
FILING DATE: 28-JUL-1994
APPLICATION NUMBER: FR 92/01064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEVEL...
CORRESPONDENCE ADDRESS:
ANDRESSEE: Rhone-Poulenc Rorer Inc.
ANDRESSEE: Anone-Poulenc Rorer Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 249 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES,
TITLE OF INVENTION: CONTAINING SAID POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: (610) 454-3839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 28-JAN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 31-JAN-1992 PRIOR APPLICATION DATA: APPLICATION NUMBER: PCT/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 500 ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Smith Ph.D., Julie K. REGISTRATION NUMBER: P-38,6 REFERENCE/DOCKET NUMBER: ST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                 APPLICANT: Fleer, Reinhard
Fournier, Alain
Guitton, Jean-Dominique
                                     NUMBER OF SEQUENCES:
                                                                                       TITLE OF
                                                                                                                                                                                                                                                                                                        232 QYSSYPWT 239
                                                                                                                                                                                                                                                                                                                                       2 QYSKLPWT 9
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                                                                                                                                                                                                                     Application US/09984186
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Yeh, Patrice
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Jung, Gerard
Yeh, Patrice
'INVENTION: NOVEL BIOLOGICALLY ACTIVE POLYPEPTIDES,
PREPARATION THEREOF AND PHARMACEUTICAL COMPOSITION
CONTAINING SAID POLYPEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (610)
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Guitton, Jean-Dominique
                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                     67.9%;
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Pred. No. 20;
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RESULT 31
$200183-5
$200183-5
;Patent No. 5200183
; APPLICANT: TANG, JORDAN J.N.;WANG, CHI-SUN
; TITLE OF INVENTION: RECOMBINANT BILE SALT J
; NUMBER OF SEQUENCES: 22
; NUMBER OF SEQUENCES: 22
; CURRENT APPLICATION DATA:
; CURRENT APPLICATION DATA:
                Query Match
Best Local Similarity
Whiches 6; Conserve
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                                                                                                       5200183-5
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FILING DATE: 31-JAN-1997
APPLICATION NUMBER: US 08/256,927
FILING DATE: 28-JUL-1994
APPLICATION NUMBER: FR 92/01064
FILING DATE: 31-JAN-1992
APPLICATION NUMBER: PCT/FR93/00085
FILING DATE: 31-JAN-1993
ATTORNEY/AGENT INFORMATION:
NAME: SMITH Ph.D., Julie K.
REGISTRATION NUMBER: P-38,619
REFERENCE/DOCKET NUMBER: ST92006-US
TELEPAX: (610) 454-3808
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                      SEQ ID NO:5:
LENGTH: 572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 75.0
                                                                                                                                                                                                                                APPLICATION NUMBER: US/
FILING DATE: 12-JUN-1990
PRIOR APPLICATION DATA:
                                                                                                                                                          APPLICATION NUMBER: 504,635
FILING DATE: 04-APR-1990
APPLICATION NUMBER: 122,410
FILING DATE: 19-NOV-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Rhone-Poulenc Rorer Inc.
STREET: 500 Arcola Road, 3C43
CITY: Collegeville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 18:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 29-Oct-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: Macintosh
OPERATING SYSTEM: System 7.1
SOPTMARE: Word 5.1 (PatentIn)
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                  232 QYSSYPWT 239
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1 HQYSKLPW 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 249 amino acids
                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       67.9%;
                                                67.9%;
75.0%;
                                                                                                                                                                                                                                                                     US/08/537,426
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Pred. No.
                                   1; Mismatches
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                                                Score 38; DB
Pred. No. 47;
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5215909-12
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Best Local Similarity '5.0
"""ches 6; Conservative
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                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446,100
FILING DATE: 19-MAY-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Hendricks, Glenna
REGISTRATION NUMBER: 32,535
REFERENCE/DOCKET NUMBER: 32,535
REFERENCE/DOCKET NUMBER: 32,535
REFERENCE/DOCKET NUMBER: 32,535
REFERENCE/DOCKET NUMBER: 57,031
TELEPHONE: (703) 425-4250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: SOREQ, HERMONA
TITLE OF INVENTION: HUMAN CHOLINESTERASE GENES
NUMBER OF SEQUENCES: 13
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/572,911
FILING DATE: 15-AUG-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 87,724
FILING DATE: 21-AUG-1987
FILING DATE: 21-AUG-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 600162
                                                                                                        TELEFAX: (703) 425-2767
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 602 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: unknown
MOLECULE TYPE: protein
HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 875,737 FILING DATE: 18-JUN-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 22031
                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Fairfax
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 573
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/08446100
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47;
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RESULT 35
US-08-446-100-3
; Sequence 3, Application US/08446100
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Best Local (
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446,100
PILING DATE: 19-MAY-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Hendricks, Glenna
REGISTRATION NUMBER: 32,535
REFERENCE/DOCKET NUMBER: broomfiel
                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (703) 425-2767 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                        TOPOLACTION TOPOLACTION TOPOLACTICAL: YES HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FRAGMENT TYPE: N-terminal ORIGINAL SOURCE: ORGANISM: human esteras
                                                                                                                                                                                                                                  FRAGMENT TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hendricks and Assoc.
                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS
LENGTH: 602 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (703) 425-4250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     451 HRSSKLPW 458
                                                                                451 HRSSKLPW 458
                                                                                                         1 HOYSKLPW 8
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                                                                                                                                        Similarity 75.06; Conservative
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                                                                                                                                                                                                                                                                                                                                                    602 amino acids
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Millard, Charles B
                                                                                                                                                                                                                                                                                                          unknown
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                                                                                                                                                        67.9%;
75.0%;
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Pred. No.
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                                                                                                                                                                   DB 3; Length 602;
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                                                                                                                                           Indels
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RESULT 36
US-08-446-100-4
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                                                                                                                                                                                                                                                                              Sequence 4, Application US/08446100 Patent No. 6001625
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Hendricks, Glenna
REGISTRATION NUMBER: 32,535
REFERENCE/DOCKET NUMBER: broo
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 425-4250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (703) 425-42:
TELEFAX: (703) 425-2767
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                            APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Millard, C. Okeana
APPLICANT: Lockridge, Okeana
TITLE OF INVENTION: Site-Directed Mutagenesis of Esterases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FRAGMENT TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446,100
FILING DATE: 19-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                        APPLICANT: Lockridge, Oksana
TITLE OF INVENTION: Site-Directed Mutagenesis of Esterases
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: unknown
MOLECULE TYPE: protein
HYPOTHETICAL: YES
ANTI-SENSE: YES
                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                    STATE: VA
COUNTRY: US
ZIP: 22031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: VA
COUNTRY: U
ZIP: 22031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 67.9%;
Local Similarity 75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                        STREET:
                                                                                                           CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: human esterases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                          ADDRESSEE:
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                                                                                                         Fairfax
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                                                                                                                    E: Hendricks and Assoc.
9669 A Main Street, P.O.
                                                                                                                                                                                                                            Broomfield, Clarence A
Millard, Charles B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Hendricks and Assoc.
9669 A Main Street, P.O. Box 2509
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Millard, Charles B
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Pred. No.
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Broomfield, Clarence A
APPLICANT: Millard, Charles B
APPLICANT: Lockridge, Oksana
TITLE OF INVENTION: Site-Directed
NUMBER OF SEQUENCES: 31
                                              TELEPHONE: (703) 425-4250
TELEPAX: (703) 425-2767
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (703) 425-42
TELEFAX: (703) 425-2767
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                             CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Hendaticks, Glenna
REGISTRATION NUMBER: 32,535
REFERENCE/DOCKET NUMBER: broo
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 425-4250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: promote yes enuge; yes
                                                                                                                                                                                              COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446,100
FILLING DATE: 19-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FRAGMENT TYPE: ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
               SEQUENCE CHARACTERISTICS:
LENGTH: 602 amino acid
                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                    CITY: Fairfax
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Hendricks, Glenna
REGISTRATION NUMBER: 32,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: sir
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         451 HRSSKLPW 458
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9669 A Main Street, P.O. Box 2509
602 amino acids
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75.0%;
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In Release #1.0, Version #1.25
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Pred. No.
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US-08-446-100-6
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                                                         Matches
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                                                                                    Query Match
                                                                                                                                                                                                                                                                           TELEFAX: (703) 425-2767 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
APPLICANT:
APPLICANT:
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ORIGINAL SOURCE:
ORGANISM: hum
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                                                                                                                                                                       MOLECULE TYPE: protein HYPOTHETICAL: YES ANTI-SENSE: YES
                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Hendricks, Glenna
REGISTRATION NUMBER: 32,535
REFERENCE/DOCKET NUMBER: br
                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 425-4250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Fairfax
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TUMBER OF SEQUENCES:
                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 19-MAY-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: si
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                                                                                                                                                                                                                                                             ENGTH:
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LE OF INVENTION: Site-Directed Mutagenesis of Esterases
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451 HRSSKLPW 458
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                         1 HOYSKLPW 8
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9669 A Main Street, P.O. Box 2509
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Millard, Charles B
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Pred. No. 50;
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Pred. No.
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                                                                                                                                                                                       US-08-446-100-8
                                                                                                                                                    Sequence 8, Application US/08446100 Patent No. 6001625
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Matches 6; Conserv
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                         GENERAL INFORMATION:
APPLICANT: Broomfield, Clarence A
APPLICANT: Milard, Charles B
APPLICANT: Lockridge, Oksana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Hendricks, Glenna
REGISTRATION NUMBER: 32,535
REFERENCE/DOCKET NUMBER: br
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                          FRAGMENT TYPE:
ORIGINAL SOURCE:
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LENGTH: 602 amino acid
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CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                            CORRESPONDENCE ADDRESS:
                                                            NUMBER OF SEQUENCES:
                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL:
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           STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: FILING DATE: 19-MAN CLASSIFICATION: 43:
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                           ADDRESSEE:
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Fairfax
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           E: Hendricks and Assoc.
9669 A Main Street, P.O. Box 2509
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9669 A Main Street, P.O. Box 2509
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Lockridge, Oksana
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                                                                         Lockridge, Oksana
VENTION: Site-Directed Mutagenesis of Esterases
                                                                                                                                                                                                                                                                                                                 Conservative
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Pred. No.
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COUNTRY: US
ZIP: 22031
COMPUTER READABLE FORM:
MEDIUM TYPE: #10ppy disk
MEDIUM TYPE: #10ppy disk
COMPUTER: ILMN CCOMPARTION:
OPERATING SYSTEM: PC-DOS/MS-DOS
CURRENT APPLICATION NUMBER: US/08/446,100
FILING DATE: 19-MA-1995
FRIEDRING PORTON: NUMBER: DATE: DAT
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Database :
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Scoring table:
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Perfect score:
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                                              Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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56
1 HQYSKLPWT 9
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Gapop 10.0 , Gapext 0.5
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185.159 Million cell updates/sec
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Copyright (c) 1993 - 2004 Compugen Ltd.
SUMMARIES
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0 0 0 0 0 0 0 4 4 4 4 4 4 4 4 4 4 4 4 4	Score
100.0 100.0 100.0 100.0 100.0 100.0 78.6 78.6 76.8 71.4 71.4	Match
128 128 128 128 128 128 109 215 215 9 9	Query Match Length DB
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Sequence 10, Appl Sequence 19, Appl Sequence 19, Appl Sequence 19, Appl Sequence 10, Appl Sequence 10, Appl Sequence 10, Appl Sequence 12, Appl Sequence 32, Appl Sequence 4290, Appl Sequence 24, Appl Sequence 26, Appl Sequence 26, Appl Sequence 27, Appl Sequence 28, Appl Sequence 5, Appl Sequence 5, Appl	Description

45	44	43	42	41	40	39 9	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14
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107	107	107	107	107	107	107	107	107	107	103	103	103	103	103	64	9	9	9	9	159	153	126	110	109	108	108	108	107	107	107	101
16	16	16	16	16	14	14	14	9	9	16	16	16	16	16	17	15	14	9	9	9	9	16	14	9	14	10	9	17	16	15	7
-10-723-	-10-723-434-	-10-723-434-	-10-723-	-10-723-434-	-10-251-215-	US-10-234-671-15	-10-234-671-1	56-160B-1	US-09-056-160B-13	-10-379-392-1	-10-379-392-	US-10-379-392-123	-10-379-392-	-10-379-392-	5-115-	0-624-153-	234-671-	US-09-217-268B-31	-160B-	US-09-187-693-66	-693-6		US-10-044-896-4	US-09-811-123-6	4-671-	US-09-795-798-3	US-09-056-160B-12	-641-1	US-10-619-754-5	-10-600-152-	-10-000-070-
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## ALIGNMENTS

CURRENT FILING DATE: 1999-01-05 EARLIER APPLICATION NUMBER: 09/225,322 EARLIER FILING DATE: 1999-01-05 EARLIER APPLICATION NUMBER: US 08/454,680 EARLIER APPLICATION NUMBER: US 08/408,133 EARLIER APPLICATION NUMBER: US 08/408,133 EARLIER FILING DATE: 1995-03-21 EARLIER FILING DATE: 1995-03-21 EARLIER FILING DATE: 1994-08-17 EARLIER APPLICATION NUMBER: US 08/292,178 EARLIER FILING DATE: 1992-09-17 EARLIER FILING DATE: 1992-09-17 EARLIER FILING DATE: 1992-09-17 EARLIER FILING DATE: 1991-09-18 NUMBER OF SEQ ID NOS: 19 SOFTWARE: Patentin Ver. 2.0 SEG ID NO 10 LENGTH: 128 TYPE: PRT ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: CDNA KM-641 US-09-764-304-10	77222220 0
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APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HANAI, NOBUO
APPLICANT: HISBGAWA, MAMORU
APPLICANT: KIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
PILE REFERENCE: 249-101
CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER APPLICATION NUMBER: US/09/64,680
EARLIER APPLICATION NUMBER: US/08/454,680
EARLIER APPLICATION NUMBER: US/08/454,680
EARLIER APPLICATION NUMBER: US/08/454,680
EARLIER FILING DATE: 1995-05-31
EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US/08/408,133
EARLIER APPLICATION NUMBER: US/08/292,178
EARLIER APPLICATION NUMBER: US/08/292,178
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CURRENT APPLICATION NUMBER: US/10/265,713
CURRENT FILING DATE: 2002-10-08
PRIOR APPLICATION NUMBER: US/09/225,322
PRIOR FILING DATE: 1999-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
                                                                                                                                                  APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUYANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
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EARLIER FILING DATE: 1992-09-17
EARLIER APPLICATION NUMBER: JP 3-238375
EARLIER FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin Ver. 2.0
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ORGANISM: Artificial Sequence
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9; Conserv
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PRIOR FILING DATE: 1994-08-17
PRIOR APPLICATION NUMBER: US07/947,674
PRIOR PILING DATE: 1992-09-17
PRIOR APPLICATION NUMBER: JP 3-238375
PRIOR FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
SOFTWARE: PATENTIN VET: 2.0
SEQ ID NO 10
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Best Local Similarity
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; OTHER INFORMATION: Description of Artificial Sequence:light chain
; OTHER INFORMATION: variable region
US-10-265-713-19
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NUMBER OF SEQ ID NOS: 19
SOFTWARE: PatentIn Ver. 2.
SEQ ID NO 19
LENGTH: 128
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Publication No
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APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAAI, HIROMASA
APPLICANT: KUWANA, YOSHHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 08/454,680 PRIOR FILING DATE: 1995-05-31 PRIOR APPLICATION NUMBER: US 08/408,133 PRIOR FILING DATE: 1995-03-21 PRIOR APPLICATION NUMBER: US 08/292,178 PRIOR FILING DATE: 1994-08-17
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PRIOR FILING DATE: 1992-09-17
PRIOR APPLICATION NUMBER: JP 3-238375
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CURRENT FILING DATE: 2002-10-08
PRIOR APPLICATION NUMBER: US/09/225,322
PRIOR FILING DATE: 1999-01-05
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PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
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ORGANISM: Artificial Sequence
PEATURE:
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Publication No. US20030166876A1
GENERAL INFORMATION:
APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 10
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OTHER INFORMATION: Description of Artificial Sequence: cDNA KM-641
:-10-166-626-10
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APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/10/166,626
CURRENT FILING DATE: 2002-06-12
PRIOR APPLICATION NUMBER: US/09/225,322B
PRIOR PILING DATE: 1999-01-05
PRIOR FILING DATE: 1999-01-05
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CURRENT APPLICATION NUMBER: US/10/166,626

CURRENT FILING DATE: 2002-06-12

PRIOR APPLICATION NUMBER: US/09/225,322B

PRIOR FILING DATE: 1999-01-05

PRIOR APPLICATION NUMBER: US 08/454,680

PRIOR FILING DATE: 1995-05-31

PRIOR FILING DATE: 1995-05-31
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Publication No. US20030166876A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: SHITARA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 128
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
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OR FILING DATE: 1995-03-21
OR APPLICATION NUMBER: US 08/292,178
OR FILING DATE: 1994-08-17
OR APPLICATION NUMBER: US07/947,674
OR APPLICATION NUMBER: US07/947,674
OR APPLICATION NUMBER: JP 3-238375
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les 9; Conserv
                                                                                                                                                                        APPLICATION NUMBER: US 08/454,680 FILING DATE: 1995-05-31 APPLICATION NUMBER: US 08/408,133 FILING DATE: 1995-03-21
                                      APPLICATION NUMBER: US 08/292,178
FILING DATE: 1994-08-17
APPLICATION NUMBER: US07/947,674
FILING DATE: 1992-09-17
APPLICATION NUMBER: JP 3-238375
FILING DATE: 1991-09-18
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PatentIn Ver. 2.0
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Best Local Similarity
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                                                              US-10-371-942-32
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US-10-697-399-6
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                                                                                          NUMBER OF SEQ ID NOS: 121
SOFTWARE: FRSTSEQ for Windows Version 4.0
SEQ ID NO 32
LENGTH: 109
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 50
SOFTWARE: PatentIn version 3.2
SEQ ID NO 6
LENGTH: 99
                Query Match
Best Local :
                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                  Sequence 32, Application US/10371942 Publication No. US20030223994A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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 Matches
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                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/371,942
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: US 60/358,994
                                                                                                                                                                                                                                                                 APPLICANT: Hoogenboom, Henricus Renerus Jacobus Mattheus
                                                                                                                                                                         PRIOR FILING DATE:
                                                                                                                                                                                                                                         FILE REFERENCE: 10280-034001
                                                                                                                                                                                                                                                    APPLICANT: Reiter, Yoram
TITLE OF INVENTION: MHC-PEPTIDE COMPLEX BINDING LIGANDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/697,399
CURRENT FILING DATE: 2003-10-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Methods of Optimizing Antibody Variable Region Binding Affinity FILE REFERENCE: AME-08122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: light chain OTHER INFORMATION: variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
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Similarity 7; Conserv
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Vasserot, Alain P.
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Smith, Eric P.
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 Conservative
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87.5%;
                78.6%;
77.8%;
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                Score 44;
Pred. No.
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Pred. No.
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Pred. No.
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 Mismatches
                               DB 14;
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                             Length 109;
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HQYGFLPWT 98

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FILE REFERENCE: P1469RIC1
CURRENT APPLICATION NUMBER: US/10/624,153
CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION NUMBER: US 09/440,781
PRIOR FILING DATE: 1999-11-16
PRIOR APPLICATION NUMBER: US 60/108,945
PRIOR FILING DATE: 1998-11-18
NUMBER OF SEQ ID NOS: 99
LENGTH: 9
LENGTH: 9
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: MISC FEATURE
LOCATION: (168)
OTHER INFORMATION: Xaa equals any of the twenty naturally occurring L-amino acids
US-10-264-049-4290
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US-10-264-049-4290
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CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/18569
PRIOR FILING DATE: 2001-06-07
PRIOR APPLICATION NUMBER: US 60/209,467
PRIOR FILING DATE: 2000-06-07
NUMBER OF SEQ ID NOS: 4360
SOFTWARE: Patentin Ver. 3.1
  Best Loc
Matches
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Publication No. US2004008502A1
GENERAL INFORMATION:
APPLICANT: CHEN, YVONNE M.
APPLICANT: LOWMAN, HENRY B.
APPLICANT: MULLER, YVES
TITLE OF INVENTION: ANTIBODY VARIANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 4290
LENGTH: 215
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Publication No. US20040005579A1
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TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA133P1
                                                                                                                      FEATURE:
NAME/KEY: artificial
LOCATION: 1-9
                                                                                                     OTHER INFORMATION: variant CDR sequence
                                                                                                                                                                                                         ORGANISM: artificial sequence FEATURE:
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                                                                                                                                                                                    OTHER INFORMATION: sequence is synthesized
  Local Similarity 75.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    76.8%; Score 43; DB 15; Local Similarity 100.0%; Pred. No. 20; nes 7; Conservative 0; Mismatches 0
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                   71.4%;
75.0%;
Score 40; DB 15; Length 9; Pred. No. 1.4e+06; 1; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-424-599-159861; Sequence 159861, Application US/10424599; Publication No. US20040031072A1
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                                                                                                                                                                                                                  APPLICANT: La Rosa Thomas J
APPLICANT: Kovalic David K
APPLICANT: Kovalic David K
APPLICANT: Cap Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21 (53223) B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEG ID NOS: 285684
SEQ ID NO 159861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/108,945
PRIOR FILING DATE: 1998-11-18
NUMBER OF SEQ ID NOS: 99
SEQ ID NO 26
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                                                                      Query Match
Best Local
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Publication No.
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APPLICANT: LOWMAN, HENRY B.
APPLICANT: MULLER, YVES
TITLE OF INVENTION: ANTIBODY VARIANTS
FILE REFERENCE: P1469R1C1
CURRENT APPLICATION NUMBER: US/10/624,153
CURRENT FILING DATE: 2003-07-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 09/440,781 PRIOR FILING DATE: 1999-11-16
                                                                                                                                                          TYPE: PRT
ORGANISM: Glycine max
FEATURE:
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NAME/KEY: artificial
LOCATION: 1-9
OTHER INFORMATION: variant CDR sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: artificial sequence
FEATURE:
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                                                                                                                                                                                                             LENGTH: 93
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6; Conserva
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6; Conserv
1 HQYSKLPWT 9
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                                                    Conservative
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                                                                      71.4%;
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75.0%;
                                                    Score 40; DB 15; Length 93; Pred. No. 29; Indels
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Pred. No.
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                 RESULT 15
US-10-600-152-14
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US-10-268-501-5
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US-10-608-626-5
Sequence 14, Application US/10600152
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Matches
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SEQ ID NO 5
LENGTH: 107
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GENERAL INFORMATION:
APPLICANT: Kelsey, Stephen M.
APPLICANT: Sliwkowski, Mark X.
TITLE OF INVENTION: Treament with Anti-ErbB2 Antibodies
                                                                                                                                                                                                     Query Match
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Best Local (
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Publication No. US20030086924A1
                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 10/268,501
PRIOR FILING DATE: 2002-10-10
PRIOR PPLICATION NUMBER: US 09/602,812
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/141,316
PRIOR FILING DATE: 1999-06-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 09/602,812
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/141,316
PRIOR FILING DATE: 1999-06-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Sliwkowski, Mark X.
TITLE OF INVENTION: Treatment with Anti-ErbB2 Antibodies
FILE REFERENCE: P1467R2P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/608,626
CURRENT FILING DATE: 2003-06-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/268,501
CURRENT FILING DATE: 2002-10-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: P1467R2P2
                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                      LENGTH: 107
                                                                                                                                                                                    Local
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                                                                                           90
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                                                                                                                              2 QYSKLPWT 9
                                                                                                                                                                   Similarity 75.
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                                                                                           QYNSLPWT 97
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75.0%;
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Pred. No. 33;
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Query Match
Best Local Similarity
Watches 6; Conserv:
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                                                                                                                                           US-10-619-754-5
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CURRENT FILING DATE: 2003-06-20
PRIOR APPLICATION NUMBER: 09/648,067
PRIOR FILING DATE: 2000-08-25
PRIOR FILING DATE: 105/0151,018
PRIOR APPLICATION NUMBER: US 60/151,018
PRIOR FILING DATE: 1999-08-27
PRIOR FILING DATE: 105/01213,822
PRIOR FILING DATE: 2000-06-23
                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 6
SOFTWARE: PREUSEQ for Windows Version 4.0
SEQ ID NO 5
LENGTH: 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 15
SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Baughman, Sharon A.
APPLICANT: Shak Steven
                                                                     Matches
                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 5, Application US/10619754 Publication No. US20040106161A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Publication No. US20040037824A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 60/480,043 PRIOR FILING DATE: 2003-06-20
                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/619,754
CURRENT FILING DATE: 2003-07-14
PRIOR APPLICATION NUMBER: US 60/396,290
PRIOR FILING DATE: 2002-07-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Genentech, Inc. APPLICANT: Koll, Hans
                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Kelsey, Stephen
TITLE OF INVENTION: Methods For Identifying Tumors That Are
TITLE OF INVENTION: Responsive To Treatment With Anti-ErbB2 Antibodies
FILE REFERENCE: 39766-0114A
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Dosages for Treatment with Anti-ErbB2 Antibodies FILE REFERENCE: P1775R1
                                                                                                                                                        TYPE: PRT
ORGANISM: Homosapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
                                                               Local Similarity tes 6; Conserv
90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           90 QYNSLPWT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                107
                               2 QYSKLPWT 9
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QYNSLPWT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bossenmaier, Birgit
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Muller, Hans-Joachim
Sliwkowski, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shak Steven
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                      75.0%;
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                                                                                                       71.48;
                                                                                      Score 40;
Pred. No.
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                                                                   Mismatches
                                                                                                       DB 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15;
                                                                                                       Length 107;
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                                                                 Indels
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RESULT 17 US-10-835-641-18

Sequence 18, Application US/10835641 Publication No. US20040236078A1

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RESULT 18
US-09-056-160B-12
Sequence 12, Application US/0
Sequence 10, US20020032315A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 71.4%;
Best Local Similarity 75.0%;
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 18: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
                                                                                                                                                                         TITLE OF INVENTION: ANTI-VEGF ANTIBODIES NUMBER OF SEQUENCES: 131
                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                  PPLICANT:
                                                                           CITY: South San Francisco
STATE: California
                                            ZIP:
                                                            COUNTRY:
                                                                                                                  STREET:
                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REGERENCE/DOCKET NUMBER: P0709P1D2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Carter, Paul J.
Presta, Leonard G.
TITLE OF INVENTION: Method for Making Humanized Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE DESCRIPTION: SEQ ID NO: 18:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                        90 QYNSLPWT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 QYSKLPWT 9
                                          94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/705,398
FILING DATE: 02-Nov-2000
APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
APPLICATION NUMBER: 07/715272
FILING DATE: 14-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/10/835,641 FILING DATE: 30-Apr-2004 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                            Application US/09056160B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 107 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: South San Francisco
                                                                                                                                                                                                        Wells, James A.
Presta, Leonard G.
Lowman, Henry B.
Chen, Yvonne M.
                                                                                                                    1 DNA Way
                                                            USA
                                                                                                                                                                                                                                                                                        Baca, Manuel
                                                                                                                                Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Mismatches
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Pred. No. 33;
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RESULT 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 3, Application US/09795798 Publication No. US20030207336A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/054,856
FILING DATE: 06-AUG-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P1093R2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 12:
                                     REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
TELEPAX: 650/922-9881
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                               COMPUTER READABLE FORM:

WEDIUM TYPE: 3.5 inch, 1.44 Mb flope

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/795,798

FILING DATE: 28-Feb-2001

CLASSIFICATION - CURROWN>

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/974,899

FILING DATE: CURROWN>

APPLICATION NUMBER: 08/974,899

FILING DATE: CURROWN>

APTORNEY/AGENT INFORMATION:

NAME: Lee, Wendy M.

REGISTRATION NUMBER: 40,378

REGISTRATION NUMBER: 0.5104481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 108 amino acids
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/056,160B FILING DATE: 06-Apr-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: I
SOFTWARE: WinPatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jardieu, Paula M.
TITLE OF INVENTION: Humanized Anti-CD11a Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Presta, Leonard G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90 QYNSLPWT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 QYSKLPWT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              71.4%;
Similarity 75.0%;
6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
LENGTH: 108 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-Apr-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                        inch, 1.44 Mb floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         P1093R2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 40;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                      P1014R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 9;
34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 108;
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RESULT 20
US-10-234-671-12
; Sequence 12, Application US/10234671
; Publication No. US20030190317A1
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RESULT 21
US-09-811-123-6
                                                                               밁
                                                                                                                                                                                                                                  ; SEQUENCE DESCRIPTION: SEQ ID NO: 12: US-10-234-671-12
                                                                                                                                                     Query Match 71. Best Local Similarity 75. Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 44,637
REFERENCE/OCKET NUMBER: P1093R2C1
TELECOMUNICATION INFORMATION:
TELEPHONE: 650/225-8674
TELEPHONE: 650/225-9881
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US/10/234,671
FILING DATE: 03-Sep-2002
CLASSIFICATION: <UNknown>
PRIOR APPLICATION DATE: 09/056160
APPLICATION NUMBER: 09/056160
FILING DATE: 06-APR-1998
APPLICATION NUMBER: 60/126446
FILING DATE: 07-APR-1997
APPLICATION NUMBER: 60/054856
FILING DATE: 07-APR-1997
APPLICATION NUMBER: 60/054856
FILING DATE: 06-AUG-1997
ATTORNEY/AGENT INFORMATION:
NAME: Col, Steven X.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                / Match 71.4%;
Local Similarity 75.0%;
nes 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: ANTI-VEGF ANTIBODIES NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Baca, Manuel
                                                                          90 QYNSLPWT 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         90 QYNSLPWT 97
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                                                                                                               2 QYSKLPWT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                           TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                           TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wells, James A. Presta, Leonard G. Lowman, Henry B. Chen, Yvonne M.
                                                                                                                                                                          71.4%;
75.0%;
                                                                                                                                                                          Score 40; DB 14; Length 108; Pred. No. 34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 40; DB
Pred. No. 34;
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RESULT 23 US-10-469-125-8

Sequence 8, Application US/10469125
Publication No. US20040143101A1
GENERAL INFORMATION:
APPLICANT: Soltis, Daniel A.
APPLICANT: Burch, Ronald M.

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APPLICANT: Chuntharapai, Anan
APPLICANT: Kim, Jin K.
APPLICANT: Kim, Jin K.
APPLICANT: Stewart, Timothy
APPLICANT: STEWART G.
FILE OF INVENTION: ANTI-INTERFERON-ALPHA ANTIBODIES
FILE REFERENCE: GENENT. 074A
CURRENT APPLICATION NUMBER: US/10/044,896
CURRENT FILING DATE: 2002-01-09
PRIOR APPLICATION NUMBER: 60/270775
PRIOR APPLICATION NUMBER: 60/270775
PRIOR FILING DATE: 2001-02-22
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
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US-10-044-896-4
                                                                                                                                                                         US-10-044-896-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/10044896 Publication No. US20030166228A1 GENERAL INFORMATION:
                                                                                Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 6
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Patent No. US20020001587A1
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Best Local (
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APPLICANT: Ralph Schwall
APPLICANT: Mark Slwkowski
TITLE OF INVENTION: METHODS OF TREATMENT USING ANTI-ErbB
TITLE OF INVENTION: ANTIBODY-WAYTANSINOID CONJUGATES
FILE REFERENCE: GENENT.073A2
CURRENT APPLICATION NUMBER: US/09/811,123
CURRENT APPLICATION 2001-03-16
CURRENT FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/238,327
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 09/602,530
PRIOR FILING DATE: 2000-06-23
NUMBER OF SEQ ID NOS: 11
                                                                                                                                                                                        LENGTH: 110
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
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2 QYSKLPWT 9
||: ||||
90 QYNSLPWT 97
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                                                                                    Conservative
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75.0%;
                                                                                                        71.4%; Score 40; 75.0%; Pred. No.
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Pred. No.
                                                                                        Mismatches
                                                                                                          DB 14;
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                                                                                                                              Length 110;
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                                                                                    Gaps
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TITLE OF INVENTION: IMMUNGLOBULIN CONSTRUCT CONTAINING ANTI-MUCIN VARIABLE DOMAIN SETTITLE OF INVENTION: FOR ELICITING AN ANTI-IDIOTYPE ANTI-TUMOR RESPONSE FILE REFERENCE: 02755/100G273-US1
CURRENT APPLICATION NUMBER: US/10/469,125
CURRENT FILING DATE: 2003-08-25
PRIOR APPLICATION NUMBER: US 60/281,182
PRIOR APPLICATION NUMBER: US 60/281,182
PRIOR APPLICATION NUMBER: PCT/US02/10304
PRIOR APPLICATION NUMBER: PCT/US02/10304
PRIOR FILING DATE: 2002-04-02
PRIOR FILING DATE: 2002-04-02
NUMBER OF SEQ ID NOS: 44
SOFTWARE: Patentin version 3.1
SEQ ID NO 8
SOFTWARE: PATENTIAL SEQUENCE                                                                                                                                      US-09-187-693-66
; Sequence 66, Application US/09187693
; Patent No. US20020173629A1
; GENERAL INFORMATION:
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US-09-187-693-62
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TYPE: PRT
ORGANISM: human
FEATURE:
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Best Local Similarity 75.0%;
Matches 6; Conservative
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Best Local S
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APPLICANT: Yang, Xiao-Dong
APPLICANT: Gallo, Michael
APPLICANT: Gallo, Michael
TITLE OF INVENTION: Human Monoclonal Antibodies to Epidermal
TITLE OF INVENTION: Growth Factor Receptor
FILE REFERENCE: Cell 4.20 CIP2
CURRENT APPLICATION NUMBER: US/09/187,693
CURRENT FILING DATE: 1998-11-05
APPLICANT: Jakobovits, Aya APPLICANT: Yang, Xiao-Dong
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LOCATION: (1)...(153)
OTHER INFORMATION: Xaa = Any Amino Acid
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Pred. No. 39;
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Pred. No. 47;
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, NAME/KEY: VARIANT
; LOCATION: (1)...(159)
; OTHER INFORMATION: Xaa = Any Amino Acid
US-09-187-693-66
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US-09-056-160B-6
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 66
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                                                                                                                     APPLICATION NUMBER: US/09/056,160B
FILING DATE: 06-Apr-1998
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 66/054,856
FILING DATE: 06-AUG-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: p1093R2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Jia, Xiao-Chi
TITLE OF INVENTION: Human Monoclonal Antibodies to Epidermal
TITLE OF INVENTION: Growth Factor Receptor
FILE REFERENCE: Cell 4.20 CIP2
CURRENT APPLICATION NUMBER: US/09/187,693
CURRENT FILING DATE: 1998-11-05
PRIOR APPLICATION NUMBER: 09/162,280
PRIOR APPLICATION NUMBER: 09/851,362
PRIOR APPLICATION NUMBER: 09/851,362
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
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APPLICANT: Baca,
APPLICANT: Wells,
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ORGANISM: human
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                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: ANTI-VEGF ANTIBODIES NUMBER OF SEQUENCES: 131
CORRESPONDENCE ADDRESS:
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Local Similarity 66.7%;
les 6; Conservative
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CITY: South San Francisco
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Jia, Xiao-Chi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genentech, Inc.
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Pred. No.
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PEATURE:

NAME/KEY: MISC FEATURE

OTHER INFORMATION: CDR of murine R3 antibody
US-09-217-268B-31
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; TOPOLOGY:
US-09-056-160B-6
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US-10-234-671-6
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Best Local Similarity 75.0%;
Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Humanized and Chimeric Monoclonal Antibodies That Recognize Epide TITLE OF INVENTION: Growth Factor Receptor (EGF-R); Diagnostic and Therapeutic Use FILE REFERENCE: 2720.10S CURRENT APPLICATION NUMBER: US/09/217,268B CURRENT FILING DATE: 1998-12-21 NUMBER OF SEQ ID NOS: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Mateo de Acosta del Rio, Christina M
APPLICANT: Rodriquez, Rolando P
APPLICANT: Frias, Ernesto M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/234,671
FILING DATE: 03-Sep-2002
                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chen, Yvonne M.
TITLE OF INVENTION: ANTI-VEGF ANTIBODIES
NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Baca, Manuel
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US20020065398A1
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                                                                                                                                                                                                                                                                                                                                                                                 CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                              ZIP: 94080
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
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75.0%;
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Pred. No. 1.4e+0
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; TOPOLOGY: Linear ; SEQUENCE DESCRIPTION: SEQ ID NO: US-10-234-671-6
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                                                                                                                                    ; FEATURE:
; NAME/KEY: artificial
; LOCATION: 1-9
; OTHER INFORMATION: variant CDR seguence
US-10-624-153-25
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                                                                   Best Loc
Matches
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PRIOR PILING DATE: 1998-11-18
NUMBER OF SEQ ID NOS: 99
SEQ ID NO 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 25, Application US/10624153 Publication No. US20040086502A1
                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: P1469R1C1
CURRENT APPLICATION NUMBER: US/10/624,153
CURRENT FILING DATE: 2003-07-21
PRIOR APPLICATION NUMBER: US 09/440,781
PRIOR FILING DATE: 1999-11-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: CHEN, YVONNE M.
APPLICANT: LOWMAN, HENRY B.
APPLICANT: MULLER, YVES
TITLE OF INVENTION: ANTIBODY VARIANTS
                                                                                                                                                                                                                          ORGANISM: artificial sequence FEATURE: OTHER INFORMATION: sequence is synthesized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                TYPE: PRT
                                                                                                                                                                                                                                                                                                 LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/126446
FILING DATE: 07-APR-1997
APPLICATION NUMBER: 60/054856
FILING DATE: 06-AUG-1997
ATTORNEY/AGENT INFORMATION:
NAME: Cui, Steven X.
REGISTRATION NUMBER: 44,637
REFERENCE/DOCKET NUMBER: 91093R2C1
                                                                                   Local
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APPLICATION NUMBER: 09/056160
FILING DATE: 06-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: 650/225-8674
                                2 QYSKLPWT 9
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                                                                   Similarity 75.0
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 QYSAVPWT 9
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                                                                                   69.6%;
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75.0%;
                                                                   Score 39; DB
Pred. No. 1.4e
1; Mismatches
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Pred. No. 1.4e+06;
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                                                                                   1.4e+06;
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RESULT 30 US-10-425-115-293070 ; Sequence 293070, Application US/10425115

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Query Match
Best Local Similarity
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SEQ ID NO 293070
LENGTH: 64
TYPE: PRT
ORGANISM: Zea mays
FEATURE:
                                                                                                                       US-10-379-392-121
                                                                                                                                                 RESULT 32
                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Humanized US-10-379-392-100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: ANTIBODY OPTIMIZATION
FILE REFERENCE: A-71386-3 463077-236
CURRENT APPLICATION NUMBER: US/10/379,392
CURRENT FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: US 60/360,843
PRIOR FILING DATE: 2002-03-01
PRIOR FILING DATE: 2002-03-01
PRIOR APPLICATION NUMBER: US 60/384,197
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2002-05-29
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                                                            Sequence 121, Application US/10379392 Publication No. US20040110226A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 184
SOFTWARE: PatentIn version 3.2
SEQ ID NO 100
LENGTH: 103
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APPLICANT: Zhou, Yihwa
APPLICANT: Zhou, Yihwa
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(5322)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
APPLICANT: Lazar, Gregory Alan
APPLICANT: Desjarlais, John Rudolf
APPLICANT: Marshall, Shannon Alicia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lazar, Gregory Alan
APPLICANT: Desjarlais, John Rudolf
APPLICANT: Marehall, Shannon Alicia
APPLICANT: Dahiyat, Bassil I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Unknown
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Clone ID: MRT4577_30360C.1.pep
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nes 6; Conserv
                                                                                                                                                                                                         90 QYSTVPWT 97
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DS20040110226A1
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75.0%;
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Pred. No.
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Pred. No. 31;
                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                             DB 16; Length 103;
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RESULT 34 US-10-379-392-129

Sequence 129, Application US/10379392
Publication No. US20040110226A1
GENERAL INFORMATION:
APPLICANT: Lazar, Gregory Alan
APPLICANT: Desjariais, John Rudolf

APPLICANT:

Desjarlais, John Rudolf Marshall, Shannon Alicia Dahiyat, Bassil I.

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NUMBER OF SEQ ID NOS: 184

SOPTWARE: PatentIn version 3.2
SEQ ID NO 123
LENGTH: 103
                                                                                                                                   ; TYPE: PRT; ORANISM: Artificial sequence; EEATURE; OTHER INFORMATION: Synthetic US-10-379-392-123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ş
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Best Local Similarity
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PRIOR FILING DATE: 2002-05-29
NUMBER OF SEQ ID NOS: 184
SOFTWARE: PatentIn version 3.2
SEQ ID NO 121
LENGTH: 103
                                                                                  Query Match
Best Local S
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Sequence 123, Application US/10379392
Publication No. US20040110226A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/379,392
CURRENT FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: US 60/360,843
PRIOR FILING DATE: 2002-03-01
PRIOR APPLICATION NUMBER: US 60/384,197
PRIOR FILING DATE: 2002-05-29
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Lazar, Gregory Alan
APPLICANT: Desjarlais, John Rudolf
APPLICANT: Marshall, Shannon Alicia
APPLICANT: Dahiyat, Bassil I.
TITLE OF INVENTION: ANTIBODY OPTIMIZATION
FILE REFERENCE: A-71386-3 463077-236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/379,392
CURRENT FILING DATE: 2003-03-03
PRIOR APPLICATION NUMBER: US 60/360,843
PRIOR FILING DATE: 2002-03-01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Dahiyat, Bassil I.
TITLE OF INVENTION: ANTIBODY OPTIMIZATION
FILE REFERENCE: A-71386-3 463077-236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
                                                                   Local Similarity
les 6; Conserv
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90 QYSTVPWT 97
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                                2 QYSKLPWT 9
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                                                                     Conservative
                                                                                    75.0%;
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75.0%;
                                                                     Score 39; DB Pred. No. 48; 1; Mismatches
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Pred. No.
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48;
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                                                                                                    Length 103;
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PRIOR APPLICATION NUMBER: US 60/360,843
PRIOR FILING DATE: 2002-03-01
PRIOR APPLICATION NUMBER: US 60/384,197
PRIOR FILING DATE: 2002-05-29
NUMBER OF SEQ ID NOS: 184
SOPTWARE: Patentin version 3.2
SEQ ID NO 129
LENGTH: 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 35
US-10-379-392-131
                                                                                                                                                                                                                                      APPLICANT: Lazar, Gregory Alan
APPLICANT: Desjarlais, John Rudolf
APPLICANT: Marshall, Shannon Alicia
APPLICANT: Dahiyat, Bassil I.
TITLE OF INVENTION: ANTIBODY OPTIMIZATION
FILE REFERENCE: A-71386-3 463077-236
CURRENT APPLICATION NUMBER: US/10/379,392
CURRENT FILING DATE: 2003-03-03
                                                                           NUMBER OF SEQ ID NOS: 184
SOFTWARE: PatentIn version 3.2
SEQ ID NO 131
LENGTH: 103
                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 131, Applic Publication No. US20 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/360,843
PRIOR FILING DATE: 2002-03-01
PRIOR APPLICATION NUMBER: US 60/384,197
PRIOR FILING DATE: 2002-05-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: ANTIBODY OPTIMIZATION FILE REFERENCE: A.71386-3 463077-236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: MISC FEATURE
LOCATION: (98). (98)
OTHER INFORMATION: Xaa at position 98 can be Phe or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Xaa at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: MISC_FEATURE LOCATION: (89)..(89)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Xaa at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: MISC_FEATURE
LOCATION: (43) ... (43)
OTHER INFORMATION: Xaa at position 43 can be Tyr,
OTHER INFORMATION: Ser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: MISC_FEATURE
LOCATION: (36) . (36)
OTHER_INFORMATION: Xaa at position 36 can be Tyr, Phe, Met or Leu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
OTHER INFORMATION: Synthetic
                                     ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: MISC FEATURE LOCATION: (87)...(87)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
                       FEATURE:
OTHER INFORMATION: Synthetic
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Local Similarity 75.0%;
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Pred. No. 48;
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Query Match
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                                                                                       US-09-056-160B-15
                                                                                                                                                                                                                                                                 US-09-056-160B-13
                                        Sequence 15, Application US/09056160B Patent No. US20020032315A1 GENERAL INFORMATION:
                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/054,856
PILING DATE: 06-AUG-1997
ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P1093R2
                                                                                                                                                                                                                                                                                                                                     TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT:
APPLICANT:
APPLICANT:
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPattin (Genentech)
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                 TYPE: Amino Acid
TOPOLOGY: I.i.
                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 107 amino acid
                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 650/225-1896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: ANTI-VEGF ANTIBODIES NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 06 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: South San Francisco STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/056,160B
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                                                                                                                                              90 QYSTVPWT 97
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Presta, Leonard
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Baca, Manuel
Wells, James A.
Presta, Leonard
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genentech, Inc.
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75.0%;
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                                                                                                                                                                                                                     Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                  P1093R2
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                    9
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RESULT 38
US-10-234-671-13
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/056,160B
FILING DATE: 06-Apr-1998
CLASSIFICATION UNMBER: 60/054,856
FILING DATE: 06-AUG-1997
AFTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E.
REGISTRATION NUMBER: 28,616
REFERENCE/DOCKET NUMBER: P1093R2
TELECOMMUNICATION INFORMATION:
TELEPAN: 650/2525-1896
TELEPAN: 650/252-1896
TELEPAN: 650/252-9881
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Best Local
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INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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APPLICANT: Chen, Yvonne M.
TITLE OF INVENTION: ANTI-VEGF ANTIBODIES
NUMBER OF SEQUENCES: 131
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 1 DNA "",
CITY: South San Francisco
STATE: California
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hes 6; Conservation
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TYPE: Amino Acid
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                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/234,671
FILING DATE: 03-Sep-2002
                                                                                                                                                                                                                                                                                         Chen, Yvonne M.
TITLE OF INVENTION: ANTI-VEGF ANTIBODIES
NUMBER OF SEQUENCES: 131
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Baca, Manuel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           90 QYSTVPWT 97
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                                                                                                                                                                                                               STATE: California
                                                                                                                                                                                                                                STREET: 1 DNA Way
CITY: South San Francisco
                                                                                                                                                                      ZIP: 94080
                                                                                                                                                                                           COUNTRY: USA
                                                                                                                                                                                                                                                                    ADDRESSEE: Genentech, Inc.
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Lowman, Henry B.
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Best Local Similarity
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US-10-234-671-15
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TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 09/056160
FILING DATE: 06-APR-1998
APPLICATION NUMBER: 60/126446
FILING DATE: 07-APR-1997
APPLICATION NUMBER: 60/054856
FILING DATE: 07-APR-1997
ATTORNEY/AGENT INFORMATION:
NAME: Cui, Steven X.
REGISTRATION NUMBER: 44,637
REFERENCE/DOCKET NUMBER: P1093R2C1
TELECOMMUNICATION INFORMATION:
                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/056160
FILING DATE: 06-APR-1998
APPLICATION NUMBER: 60/126446
FILING DATE: 07-APR-1997
APPLICATION NUMBER: 60/054856
FILING DATE: 06-AUG-1997
                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                             ATTORNEY/AGENT INFORMATION:
NAME: Cui, Steven X.
REGISTRATION NUMBER: 44,637
                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/234,671
FILING DATE: 03-Sep-2002
CLASSIFICATION: Unknown>
    REFERENCE/DOCKET NUMBER: P1093R2C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-8674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: ANTI-VEGF ANTIBODIES NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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CITY: South San F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: Amino Acid TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 107 amino acids
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Lowman, Henry B.
Chen, Yvonne M.
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Pred. No. 5

    Mismatches

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US-10-251-215-49

Sequence 49, Application US/10251215

Publication No. US20030219839A1

GENERAL INFORMATION:

APPLICANT: Bowdish, Katherine S.
APPLICANT: Frederickson, Shana

TITLE OF INVENTION: ANTI-POGF ANTIBODIES AND METHODS FOR PRODUCING ENGINEERED

TITLE OF INVENTION: ANTI-POGF ANTIBODIES AND METHODS FOR PRODUCING ENGINEERED

CURRENT FILING DATE: 2002-09-20

PRIOR APPLICATION NUMBER: US/10/251,215

CURRENT FILING DATE: 2001-09-20

PRIOR APPLICATION NUMBER: US 60/333,537

PRIOR FILING DATE: 2001-09-20

PRIOR APPLICATION NUMBER: US 60/323,544

PRIOR APPLICATION NUMBER: US 60/323,544

PRIOR APPLICATION NUMBER: US 60/323,544

PRIOR RILING DATE: 2001-09-20

PRIOR RILING DATE: 2002-05-13

NUMBER OF SEQ ID NOS: 79

SOFTWARE: Patentin version 3.2

LENGTH: 107

TYPE: PRT

ORGANISM: artificial sequence

PEATURE:

OTHER INFORMATION: Humanized Light Chain

US-10-251-215-49
Search completed: December 17, 2004, 18:38:02 Job time : 18.3933 secs
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INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 107 amino acids
TYPE: Amino Acid
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GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.

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Title:
Perfect score:
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Query Matc Best Local Matches	RESULT 2 E39740 Ig kappa chain V region (Py54) C;Species: Mus musculus (house C;Date: 18-Oct-1991 #sequence_r C;Accession: E38740 R;Ruff-Jamison, S; Campos-Gonz J. Biol. Chem. 266, 6607-6613, A;Fittle: Heavy and 11ght chain A;Reference number: A38740; MUI A;Accession: E38740 A;Status: preliminary; nucleic A;Molecule type: mRNA A;Residues: 1-111 <rup> A;Cross-references: UNIPROT:091 C;Supwords: heterotetramer; imm F;19-93/Domain: immunoglobulin</rup>	Query Matches Best Local Matches Qy Qy Db 9:	30 35 62.5 125 2 31 35 62.5 140 2 32 35 62.5 318 2 33 35 62.5 318 2 34 35 62.5 318 2 34 35 62.5 318 2 35 62.5 413 2 36 35 62.5 440 2 37 35 62.5 492 2 39 35 62.5 492 2 40 35 62.5 549 2 41 35 62.5 549 2 42 35 62.5 549 2 43 35 62.5 549 2 44 35 62.5 549 2 42 35 62.5 549 2 43 35 62.5 549 2 44 35 62.5 102 2 45 35 62.5 574 2 46 35 62.5 58 2 47 35 62.5 102 2 48 35 62.5 102 2 49 35 62.5 102 2 40 35 62.5 102 2 41 35 62.5 102 2 42 35 62.5 102 2 43 35 62.5 102 2 44 35 62.5 102 2 45 35 62.5 102 2 46 6607 6613, 19 47 110: Heavy and light chain varieties Heavy and light chain variet
h Similarity 8; Conserva	Residues: 1-111 -RUF- Tresidues: 1-111 -RUF- Recidues: 1-111 -RUF- Ruff-Jamison, S.; Campos-Gonzale Biol. Chem. 266, 6607-6613, 199 Title: Heavy and 11ght chain var Reference number: A38740; MUID:9 Accession: E38740 Status: preliminary; nucleic aci. Molecule type: mRNA Residues: 1-111 -RUF- Cross-references: UNIPROT:091WS9 Superfamily: immunoglobulin V re- Reywords: heterotetramer; immuno	similarity 8; Conserv 2 QYSKLPWT 9	35 62.5 35 62.5 36 62.5 37 62.5 38 62.5 38 62.5 38 62.5 39 62.5 30 62.5 30 62.5 31 62.5 32 62.5 33 62.5 35 62.5 36 62.5 37 62.5 38 740 60 78 78 78 78 78 78 78 78 78 78 78 78 78
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R;Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R. J. Biol. Chem. 266, 6607-6613, 1991
A;Title: Heavy and light chain variable region sequences and A;Reference number: A38740; MUID:91177923; PMID:1706720
A;Accession: C38740
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C;Superfamily: immunoglobulin V region; imm
C;Keywords: heterotetramer; immunoglobulin
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A;Title: Heavy and light chain variable region sequences and A;Reference number: A38740; MUID:91177923; PMID:1706720
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Ig kappa chain V-I region (ISE) - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 19-Dec-1993 #sequence revision 18-Nov-1994 #text_change 21-Jan-2000
C;Accession: A49134; S25115
R;Rocca, A.; Khamlichi, A.A.; Aucouturier, P.; Noel, L.H.; Denoroy, L.; Preud'homme, J.L.
Clin. Exp. Immunol. 91, 506-509, 1993
A;Title: Primary structure of a variable region of the V kappa I subgroup (ISE) in light A;Reference number: A49134; MUID:93185310; PMID:7680298
A;Accession: A49134
                                                                                                                              A;Cross-references: GDB:136264
A;Map position: 2p12-2p12
A;Map position: heterotetramer subunit consists of two C;Complex: An immunoglobulin heterotetramer subunit consists of two hain disulfide bonds. In some cases, such as IgA and IgM, the subunic;Superfamily: immunoglobulin V region; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Residues: 1-128 <GAI>
A;Experimental source: hybridoma 6G6.C4
A;Experimental source: hybridoma 6G6.C4
A;Note: sequence extracted from NCBI backbone (NCBIN:134419, NCBIP:134420)
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;36-110/Domain: immunoglobulin homology <IMM>
                                                               C;Keywords: heterotetramer F;16-90/Domain: immunoglobulin homology <IMM>F;23-88/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                                                                                                                R;Goni, F.; Frangione, B.
Proc. Natl. Acad. Sci. U.S.A. 80, 4837-4841, 1983
A;Title: Amino acid sequence of the Fv region of a human
A;Reference number: A93964; MUID:83273707; PMID:6410398
A;Accession: A01876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross references: EMBL:X67322; NID:g33268; PIDN:CAA47736.1; PID:g33269
A;Note: sequence extracted from NCBI backbone (NCBIP:127088)
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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A49134
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C;Date: 03-Aug-1984 #sequence_revision 03-Aug-1984 #text_change
                                                                                                                                                                                                                                               A; Gene: GDB: IGKV1
                                                                                                                                                                                                                                                                      C; Genetics:
                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPROT:P01610 C;Comment: This chain was obtained
                                                                                                                                                                                                                                                                                                                                          A;Residues:
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Ig kappa chain V-I region (WEA) -
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                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: protein
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Best Local Similarity
Matches 6; Conserv
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Best Local
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Pred. No. 1.8;
0; Mismatches
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Pred. No.
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  39;
No.
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C;Date: U3-LC
C;Accession: T39701
R;Wood, V.; Skelton
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A;Residues: 1-258 <KUR>
A;Cross-references: UNIPROT:Q8YSA0; GB:BA000019;
A;Experimental source: strain PCC 7120
C;Genetics:
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C;Species: Nostoc sp. PCC 7120
A;Note: Nostoc sp. strain PCC 7120 is a synonym of Anabaena sp. strain PCC 7120
C;Date: 14-Dec-2001 #sequence_revision 14-Dec-2001 #text_change 09-Jul-2004
C;Accession: AF2204
C;Accession: AF2204
R;Kaneko, T.; Nakamura, Y.; Wolk, C.P.; Kuritz, T.; Sasamoto, S.; Watanabe, A.; Iriguchi Nakazaki, N.; Shimpo, S.; Sugimoto, M.; Takazawa, M.; Yamada, M.; Yasuda, M.; Tabata, S DNA Res. 8, 205-213, 2001
A;Reference number: AB1807; MUID:21595285; PMID:11759840
A;Accession: AF2204
A;Accession: AF2204
A;Accession: AF2204
                                                                                                                RESULT 10
T39701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 probable phosphotransferase - Streptomyces coelicolor C;Species: Streptomyces coelicolor C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004 C;Accession: T36237
                                                Lipoate-protein ligase A - fission yeast (Schizosaccharomyces pombe)
C;Species: Schizosaccharomyces pombe
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 09-Jul-2004
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A;Cross-references: UNIPROT:Q9X8F0;
A;Experimental source: strain A3(2)
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A;Accession: T36237
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C; Superfamily:
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V.; Skelton, J.; Churcher, C.M.; Rajandream, M.A.; ed to the EMBL Data Library, July 1999
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Pred. No.
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Pred. No.
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                     Barrell, B.G.
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   A;Molecule type: DNA
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N; Alternate names: butyrylcholinesterase
C; Species: Oryctolagus cuniculus (domestic rabbit)
C; Species: Oryctolagus cuniculus (domestic rabbit)
C; Date: 14-Feb-1992 #sequence_revision 01-Mar-1996 #text_change 09-Jul-2004
C; Accession: S10255; C39768
C; Accession: S10255; C39768
R; Jbilo, O.; Chatonnet, A.
Nucleic Acids Res. 18, 3990, 1990
A; Title: Complete sequence of rabbit butyrylcholinesterase.
A; Reference number: S10255; MUID:90326526; PMID:2374720
C;Species: Homo sapiens (man)
C;Date: 30-Jun-1987 #sequence revision 23-Feb-1996 #text_change 09-Jul-2004
C;Accession: A33769; A26613; Ā33887; A34668; A00772
R;Arpagaus, M.; Kott, M.; Vatsis, K.P.; Bartels, C.F.; La Du, B.N.; Lockrid Biochemistry 29, 124-131, 1990
A;Title: Structure of the gene for human butyrylcholinesterase. Evidence fo A;Reference number: A33769; MUID:90212557; PMID:2322535
A;Accession: A33769
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A;Title: Use of the polymerase chain reaction for homology probing of butyrylcholinester: A;Reference number: A39768; MUID:91201348; PMID:2016308
A;Accession: C39768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Introns: 485/2; 541/1
C;Superfamily: cholinest
C;Keywords: carboxylic (
F;35-535/Domain: choline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Reference number: Z21870
A;Accession: T39701
A;Status: preliminary; tran
                                                                                                                                                                                                 cholinesterase (EC 3.1.1.8) precursor [validated] - human N;Alternate names: acylcholine acylhydrolase; butyrylcholinesterase; choline
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: GB:M62779; NID:g164788; PIDN:AAA31169.1; PID:g164789
C;Genetics:
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A; Residues: 75-215 <ARP>
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A;Experimental source: strain 972h-; cosmid c17A3
C;Genetics:
A;Gene: p1038
                                                                                                                                                                                                                                                                              RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Accession: S10255
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
6; Conserve
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5; Conserv
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Pred. No. 45;
1; Mismatches
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Pred. No.
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45;
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PID:G19158

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R;Taylor, P.
submitted to the EMBL Data
                                                                                                                                                                         cholinesterase (EC 3.1.1.8) - mouse
N;Alternate names: butyryLcholine esterase
C;Species: Mus musculus (house mouse)
C;Date: 28-Oct-1996 #sequence revision 08-1
C;Accession: S70849; S15680; A39768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: DNA
A; Molecule type: DNA
A; Residues: 143-145, 'VSNWNIIFTCL' <NOG>
A; Residues: 143-meshift mutant in codon for residue 145 (Gly)
A; Lockridge, O.; Bartels, C.F.; Vaughan, T.A.; Wong, C.K.; Norton, S
J. Biol. Chem. 262, 549-557, 1987
A; Title: Complete amino acid sequence of human serum cholinesterase.
A; Reference number: A00772; MUID:87109144; PMID:3542989
A; Accession: A00772
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A;Cross-references: UNIPROT: P06276; GB::M32391; GB::J02879
A;Note: two ATG codons found upstream of Met-1 do not lie in a favorable c
R;Prody, C.A.; Zevin-Sonkin, D.; Gnatt, A.; Goldberg, O.; Soreq, H.
Proc. Natl. Acad. Sci. U.S.A. 84, 3555-3559, 1987
A;Title: Isolation and characterization of full-length cDNA clones coding
A;Reference number: A26613; MUID:87231856; PMID:3035536
A;Accession: A26613
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A;Residues: 'MSVQSNLQAGAAAASCISPKYYMIFTPCKLYHLCCRESEIN',1-602 <MCT>
A;Residues: 'MSVQSNLQAGAAAASCISPKYYMIFTPCKLYHLCCRESEIN',1-602 <MCT>
A;Rote: two ATG codons found upstream of Met-1 do not lie in a favorable context for t
A;Nogueira, C.P.; McGuire, M.C.; Graeser, C.; Bartels, C.F.; Arpagaus, M.; Van der Spe
Am. J. Hum. Genet. 46, 934-942, 1990
A;Title: Identification of a frameshift mutation responsible for the silent phenotype
A;Reference number: A34668; MUID:90252779; PMID:2339692
A;Residues:
   A; Molecule type: nucleic acid A; Residues: 1-603 < TAY>
                                                          A; Reference number: A; Accession: S70849
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Description: hydrolyzes acylcholines to choline and a carboxylic & A;Note: this cholinesterase is highly reactive with organophosphate C;Superfamily: cholinesterase; cholinesterase homology C;Keywords: carboxylic ester hydrolase; glycoprotein; homotetramer
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C;Function:
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A;Map position: 3q26.1-3q26.2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: mRNA
A; Residues: 1-133, 'D', 135-602 < PRO>
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Pred. No.
1; Mismatc
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R;Arpagaus, M.; Chatonnet, A.; Masson, P.; Newton, M.; Vaughan, T.A.; Bartels, C.F.; Nog. J. Biol. Chem. 266, 6966-6974, 1991
A;Title: Use of the polymerase chain reaction for homology probing of butyrylcholinesters A;Reference number: A39768; MUID:91201348; PMID:2016308
A;Accession: A39768
                                                                                                                                                                                                                                                       A; Map position: segment L2
C; Superfamily: reovirus mR
C; Keywords: core protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mRNA guanylyltransferase (EC 2.7.7.50) - reovirus N;Alternate names: lambda 2 protein; mRNA capping C;Species: reovirus type 3
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C;Superfamily: cholinesterase; cholinesterase homology
C;Keywords: carboxylic ester hydrolase; 9lycoprotein
F;57-557/Domain: cholinesterase homology <CHE>
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A;Molecule type: DNA
A;Residues: 97-128,'P',130-237 <ARP>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Neuron 5, 317-327, 1990
A;Title: Molecular cloning of mouse acetylcholinesterase:
A;Reference number: JH0314; MUID:90380429; PMID:2400605
A;Accession: S15680
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RESULT
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J. Biol. Chem. 262, 16289-16293, 1987
A;Title: Complete nucleotide sequence of re
A;Reference number: A28471; MUID:88058999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Note: host Homo sapiens (man)
C;Date: 30-Sep-1989 #sequence_revision 30-Sep-1989 #text_change
C;Accession: A28471
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A;Molecule type: DNA
A;Residues: 1-1289 <SEL>
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9; PMID:2824487
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Ig kappa chain V.I region (Au) - human (Species: Homo sapiens (man) C;Date: 24-Apr.1984 #sequence revision C;Accession: A91653; A01862; S02573 R;Schiechl, H.; Hilschmann, N. Hoppe-Seyler's Z. Physiol. Chem. 353, 3, 7itle: Die Primaerstruktur einer mond

A; Reference number: A91653; MUID: 72189444;

53, 345-370, 1972 monoklonalen Immunglobulin-L-Kette 2189444; PMID:5028201

vom kappa-Typ,

Subs

02-Jul-1998 #text\_change

09-Jul-2004

protein

A;Accession: A91653

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A;Note: the sequence was determined from the differentiated gene
A;Note: the authors translated the codon TGC for residue 76 as Trp
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light hain disulfide bonds. In some cases, such as IGA and IGM, the subunits associate int C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: signal sequence #status predicted <SIG>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Species: Homo sapiens (man)
C;Date: 30-Jun-1987 #sequence_revision
C;Accession: A01905
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K4HU17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Marsh, P.; Mills, F.; Gould, H.
Nucleic Acids Res. 13, 6531-6544, 1985
A;Title: Detection of a unique human VkappaIV germline :
A;Reference number: A01905; MUID:86041854; PMID:2997713
A;Accession: A01905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Complex: an immunoglobulin heterotetramer subunit consists of two identical light (kap hain disulfide bonds; in some cases, such as IgA and IgM, the subunits associate into la C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin C;Keywords: heterotetramer; immunoglobulin F;16-90/Domain: immunoglobulin homology <IMM>
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A;Title: The structure determination of the variable portion
A;Reference number: A90729; MUID:77022433; PMID:1234024
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A; Residues: 1-134 < MAR>
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                                                                                                                                                                                                    ;115-121/Region: complementarity-determining;122-134/Region: framework 4
                                                                                                                                                                                                                                                                                         ;44-60/Region: complementarity-determining;61-75/Region: framework 2
                                                                                                                                                                                                                                                                                                                                                          ;21-134/Product: Ig kappa chain V-IV region (B17);21-43/Region: framework 1
                                                                                                                                                                                                                                                                    ;76-82/Region: complementarity-determining
                                                                                                                                                                                                                                                                                                                                   ;36-116/Domain: immunoglobulin homology <IMM;
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;Comment: This is a Bence Jones protein.
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                                                                                                                                                                             43-114/Disulfide
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                                                                                                                                     Query Match
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Pred. No.
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S47329
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T31291
C;Species: Saccharomyces cerevisiae
C;Date: 02-Dec-1994 #sequence_revision 02-Dec-1994 #text_change 09-Jul-2004
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Ig kappa chain precursor V region - human (Species: Homo sapiens (man) (C;Species: Homo sapiens (man) (C;Date: 19-Dec-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 (C;Accession: A49137; S24479 (C;Accession: A49137; S24479 (C;Khamlichi, A.A.; Aucouturier, P.; Silvain, C.; Bauwens, M.; Touchard, G.; Preud'homme, Clin. Exp. Immunol. 87, 122-126, 1992 (A;Title: Primary structure of a monoclonal kappa chain in myeloma with light chain deposs A;Fattle: primary structure of a monoclonal kappa chain in myeloma with light chain deposs A;Fattle: A;Fattle: A49137; MUID:92127887; PMID:1733627
OXA1 protein precursor - yeast (Saccharomyces N;Alternate names: PET1402 protein; protein YEC;Species: Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein 1251 - Sphingomonas aromaticivorans plasmid pNL1 C;Species: Sphingomonas aromaticivorans C;Date: 11-Jan-2000 #sequence_revision 11-Jan-2000 #text_change 09-Jul-2004 C;Accession: T31291
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A;Description: Complete sequence of a 184 kb catabolic plasmid from Sphingomonas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       submitted to the EMBL Data Library, January 1992
A;Reference number: S24479
A;Accession: S24479
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A;Note: sequence extracted from NCBI backbone (NCBIN:79324; NCBIP:79325)
R;Cogne, M.C.C.
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Best Local
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A;Cross-references: GB:X07558
R;Bauer, M.; Behrens, M.; Esser, K.; Michaelis, G.; Pratje, E.
Mol. Gen. Genet. 245, 272-278, 1994
A;Title: PET1402, a nuclear gene required for proteolytic processing A;Reference number: S51575; MUID:95115677; PMID:7816036
A;Accession: S51575
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A;Title: OXAl, a Saccharomyces cerevisiae nuclear gene whose sequence is A;Reference number: S46384; MUID:94254098; PMID:8196054
A;Accession: S46384
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A;Title: Divergent overlapping transcripts at the PET122 locus in Saccharomyces cerevisi A;Reference number: A36328; MUID:90258894; PMID:2160592
A;Accession: A36328
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A; Molecule type: DNA
A; Residues: 1-585 < N
                                                                    A; Reference number: A; Accession: T47364
                                                                                                             Mayer, K.F.X.
submitted to the Protein
                                                                                                                                                           C; Accession: T47364
R; Nyakatura, G.; Fartmann,
                                                                                                                                                                                                       hypothetical protein F7M19.70 - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 09-Jul-2004
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A;Map position: 5R
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A; Residues: 1-402 < BAU>
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A; Residues: 1-379 < OHM >
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A; Residues: 1-107, 'R', 109-402 <BON>
A; Cross-references: EMBL: X77558
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A; Residues: 1-402 < DIE>
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A; Accession: S50657
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A;Description: The sequence of S. cerevisiae cosmids 8229, 9115, 9132, 9981, and lambda
                                              A; Status: preliminary
                                                                                                                                                                                                                                                                                     T47364
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
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                                                                                                                                                                                                                                                                                                                                                                               HVYSGLPW 128
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                                                                                                                  Sequence Database,
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75.0%;
                                                                                                                                                                В.;
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                                                                                                                                                                  Dauner,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 37;
Pred. No.
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47;
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R;Read, T.D.; Brunham, R.C.; Shen, C.; Gill, S.R.; Heidelberg, J.F.; White, O.; Hicker, C.; Dodson, R.; Gwinn, M.; Nelson, W.; DeBoy, R.; Kolonay, J.; McClarty, G.; Salzbern Nucleic Acids Res. 28, 1397-1406, 2000

A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae AR39

A;Reference number: A81500; MUID:20150255; PMID:10684935
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H71475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  conserved hypothetical protein TC0120 [imported] - Chlamydia muridarum (strain Nigg) C;Species: Chlamydia muridarum, Chlamydia trachomatis MoPn C;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 09-Jul-2004 C;Accession: C81739
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C81739
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Title: Genome sequence of an obligate intracellular pathogen of humans: Chlamydia track
A;Reference number: A71570; MUID:99000809; PMID:9784136
A;Accession: H71475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           probable chltr phosphoprotein - Chlamydia trachomatis (serotype D, strain UC;Species: Chlamydia trachomatis (C;Species: Chlamydia trachomatis C;Date: 13-Sep_1998 #sequence_revision 13-Sep-1998 #text_change 09-Jul-2004 C;Date: 13-Sep_1998 #sequence_revision 13-Sep_1998 #text_change 09-Jul-2004 C;Date: 14-Sep_1998 #sequence_revision 13-Sep_1998 #sequence_revision 14-Sep_1998 #sequence_revision 14-Sep_1
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A; Map position: 3
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A;Experimental source: cultivar Columbia; BAC clo
C;Genetics:
                                                                                                                                                                                                                     A; Experimental C; Genetics:
                                                                                                                                                                                                                                                   A;Molecule type: DNA
A;Residues: 1-823 <TET>
A;Cross-references: UNIPROT:09PLIO; GB:AE002279; GB:AE002160; NID:g7190148; PIDN:AAF38998
A;Experimental source: strain Nigg (MoPn)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Status: preliminary A;Molecule type: DNA
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Science 282, 754-759, 1998
                                                                                                                                                                                           A;Gene:
                                                                                                                                                                                                                                                                                                                                                                               A;Status: preliminary
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Gene: TC0120
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6; Conserv
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Pred. No.
                                                                                           Score 37;
Pred. No.
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Pred. No.
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272 HALSKLPW

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Burronylurea receptor - golden hamster
C;Species: Mesocricetus auratus (golden hamster)
C;Date: 05-May-1995 #sequence_revision 05-May-1995 #text_change 02-Feb-2001
C;Accession: A56248
                                                                                                                                                  RESULT 25
KVMS13
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A;Cross-references: GB:L40623
C;Superfamily: unassigned ATP
C;Keywords: ATP; glycoprotein
F;2-1582/Product: sulfonylure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: EMBL:X54753; NID:g55294; PIDN:CAA38555.1; PID:g55295
A;Experimental source: spleen, strain BALBC-nu/nu
A;Experimental source: spleen, strain BALBC-nu/nu
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1990
A;Note: An immunoglobulin heterotetramer subunit consists of two identical light (kag hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into labain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into labain disulfide bonds.
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A;Title: Structure of extrachromosomal circular DNAs generated by immunoglobulin light A;Reference number: S18731; MUID:91209891; PMID:1902191
A;Accession: S18731
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                            Ig kappa chain V region (PC2413) - mouse (;Species: Mus musculus (house mouse) (;Species: Mus musculus (house mouse) (;Date: 01-Sep-1981 #sequence_revision 01-Sep-1981 #text_change 09-Jul-2004 C;Accession: A01932
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A;Molecule type: DNA
A;Residues: 1-103 <HIR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Accession: S18731
R;Hirama, T.; Takeshita, S.; Yoshida, Y.; Yamagishi, H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ig kappa chain V-J region (MSI-N17) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 05-Dec-1998 #text_change 23-Jul-1999
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A;Reference number: A56248; MUID:95232532; PMID:7716547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Aguilar-Bryan, L.; Nichols, C.G.; Wechsler, S.W.; Clement IV, J.P.; Boyd III, Science 268, 423-426, 1995
R;Weigert, M.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;1362-1555/Domain: ATP-binding cassette homology <ABC2>
;1379-1386/Region: nucleotide-binding motif A (P-loop)
;10/Binding site: carbohydrate (Asn) (covalent) #status experimental
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Gatmaitan, L.; Loh, E.; Schilling, J.; Hood, L.
                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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Pred. No.
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Ryfheologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; Wh Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, ansen, N.F.; Hughes, B.; Huizar, L.
Nature 408, 816-820, 2000
A; Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khay C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Mait Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.
A; Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.
A; Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
A; Reference number: A86141; MUID:21016719; PMID:11130712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Title: Rearrangement of genetic information may produce immunoglobulin diversity.
A;Reference number: A93204; MUID:79073152; PMID:103003
A;Recession: A01932
A;Molecule type: protein
A;Residues: 1-111 <WEI>
A;Cross-references: UNIPROT:P01657
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kapphain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into lard;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer
F;16-94/Domain: immunoglobulin homology <IMM>
F;23-92/Disulfide bonds: #status predicted
                                                                                                                                           hypothetical protein W05B10.4 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #te: C;Accession: T26193
R;Mortimore, B.
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-588 <WIL>
                                                                        submitted to the EMBL Data Library, A; Reference number: Z20168
A; Accession: T26193
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T26193
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C;Superfamily: Escherichia coli trypsin-like proteinase degS;
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A;Molecule type: DNA
A;Residues: 1-518 <STO>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein F1E22.2 [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
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Best Local S
Matches 5
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Best Local (
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                                                                                                                                                                                              #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
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85.7%;
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Pred. No.
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Pred. No.
                                                                                                                         July 1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 518
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Dewar, K.;
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A;Cross-references: UNIPROT:Q23176; EMBL:Z77670; PIDN:CAB01250.1; GSPDB:GN00023; CESP:W0

HQHYSTPWT 102

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A; Experimental C; Genetical
                                                                                                                                                                                                                                   Ig light chain V region - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1998 #sequence_revision 19-Mar-1998 #text_change 21-Jan-2000
C;Accession: S26336
R;Stark, S.E.; Caton, A.J.
J. Exp. Med. 174, 613-624, 1991
A;Title: Antibodies that are specific for a single amino acid interchange in A;Reference number: S26309; MUID:91341421; PMID:1908510
A;Accession: S26336
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S26336
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F;35-49/Region: framework 2
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R;Shlomchik, M.; Mascelli, M.; Shan, H.; Radic, M.Z.; Pisetsky, D.; Marshak-Rothstein, A J. Exp. Med. 171, 265-297, 1990

A;Title: Anti-DNA antibodies from autoimmune mice arise by clonal expansion and somatic A;Reference number: PL0231; MUID:90111618; PMID:2104919

A;Accession: PL0262
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A; Residues: 1-109 < STA>
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F;57-88/Region: framework 3
F;89-97/Region: complementarity-determining 3
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A; Residues: 1-106 <SH
C; Superfamily: immuno
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A; Introns: 25/3;
C; Superfamily: k
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                                                   Query Match
Best Local S
Matches 5
                                                                                                                        Superfamily: immunoglobulin V region; immunoglobulin homology;Keywords: heterotetramer; immunoglobulin;15-95/Domain: immunoglobulin homology <IMM>
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;Species: Mus musculus (house mouse)
;Date: 16-Sep-1992 #sequence_revision 16-Sep-1992 #text_change 21-Jan-2000
                                                                                                                                                                                Cross-references: EMBL:X59201
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Superfamily: human alpha,alpha-trehalase
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Best Local
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HQYSKLPWT 9
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Pred. No. 27;
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Pred. No. 2
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Pred. No.
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                                                                                      Length 109;
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RESULT 32

B87293
hydrolase, probable [imported] - Caulobacter crescentus
c;Species: Caulobacter crescentus
c;Species: Caulobacter crescentus
c;Apecies: Caulobacter crescentus
c;Apecies: 20-Apr-2001 #sequence_revision 20-Apr-2001 #text_change 12-Jul-2004
c;Accession: B87293
R;Nierman, W.C.; Feldblyum, T.V.; Paulsen, I.T.; Nelson, K.E.; Eisen, J.; Heidelberg, J.;
B.; Laub, M.T.; DeBoy, R.T.; Dodson, R.J.; Durkin, A.S.; Gwinn, M.L.; Haft, D.H.; Kolona
n, J.; Ermolaeva, M.; White, O.; Salzberg, S.L.; Shapiro, L.; Venter, J.C.; Fraser, C.M.
Proc. Natl. Acad. Sci. U.S.A. 98, 4136-4141, 2001
A;Title: Complete Genome Sequence of Caulobacter crescentus.
A;Reference number: A87249; MUID:21173698; PMID:11259647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Kaine, B.P.; Sykes, Smith, H.O.; Woese, C.R.; Venter, J.C. A;Title: The complete genome sequence of the hyperthermophilic, sulfate-reducing archa A;Reference number: A69250, MUID:98049343; PMID:9389475
A;Accession: H69405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Residues: 1-125 <KLE>
A;Ross-references: EMBL:X72443; NID:g441354; PIDN:CAA51111.1; PID:g441355
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;34-108/Domain: immunoglobulin homology <IMM>
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$4033

Ig kappa chain V-J region - human

Ig kappa chain V-J region - human

C;Species: Homo sapiens (man)

C;Date: 19-May-1994 #sequence_revision 26-May-1995

C;Accession: $40333

C;Accession: $40333

C;Accession: $40333

C;Accession: $40333
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eur. J. Immunol. 23, 3248-3271, 1993
A;Title: Expressed human immunoglobulin chi genes and their A;Reference number: S40312; MUID:94080891; PMID:8258341
A;Accession: S40333
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-140 <KLE>
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Best Local S
Matches 6
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Best Local S
Matches 5
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Pred. No.
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Pred. No.
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al protein AF1249
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EYSSLPW

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A;Molecule 'Y'...
A;Residues: 1-312 <WIL>
A;Cross-references: UNIPROT:Q9U2A0;
A;Cross-references: clone Y48B6A
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                                                                                                                                                                                                                     A;Title: Complete genome sequence of Pseudomonas aeruginosa
A;Reference number: A82950; MUID:20437337; PMID:10984043
A;Accession: B83527
                                                                                                                                                                                                                                                                       Nature 406, 959-964, 2000
                                                                                                                                                                                                                                                                                                     C;Accession: B83527
R;Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypothetical protein Y48B6A.14 - C;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_rev
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A;Molecule type: DNA
A;Residues: 1-296 <STO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Accession: B87293
                                                                                                                                                                                                                                                                                                                                                                                        hypothetical
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                                                                                                                                           A;Cross-references: UNIPROT:Q9I503; GB:AE004529; GB:AE004091; NID:g9946851; PIDN:AAG043·A;Experimental source: strain PAO1
                                                                                                                                                                                            A; Molecule type: DNA
                                                                                                                                                                                                       A; Status: preliminary
                                                                                                               ;Gene: PA0955
                                                                                                                                                                             ;Residues: 1-318 <STO>
                                                                                                                                                                                                                                                                                                                                              ;Species: Pseudomonas aeruginosa
;Date: 15-Sep_2000 #sequence_revision 15-Sep-2000 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;Gene: CESP:Y48B6A.14
;Introns: 132/1; 210/1;
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                                                                Query Match
Best Local
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Best Local
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les 5; Conserv
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                  N
                                              Similarity 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                     protein PA0955 [imported] -
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2 QYSKILPW
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                                                                62.5%;
71.4%;
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83.3%;
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71.4%;
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                                                1;
                                                                Score 35;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           September 1999
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                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                      Pseudomonas aeruginosa (strain PAO1)
                                                                84
                                                                              DB 2;
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                                                                              Length 318;
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                                                Indels
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                                                Gaps
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                                                                                                                                                                                                                                                                                                        M.J.; B:
K.; Lim,
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hypothetical protein [imported] - Staphylococcus aureus (strain N315) (;Species: Staphylococcus aureus (;Species: Staphylococcus aureus C;Date: 10-May-2001 #sequence_revision 10-May-2001 #text_change 09-Jul-2004 C;Accession: A89761 R;Kuroda, M.; Ohta, T.; Uchiyama, I.; Baba, T.; Yuzawa, H.; Kobayashi, I.; Cui, ma, A.; Mizutani-Ui, Y.; Kobayashi, N.; Sawano, T.; Inoue, R.; Kaito, C.; Sekimi C.; Shiba, T.; Hattori, M.; Ogasawara, N.; Hayashi, H.; Hiramatsu, K. Lancet 357, 1225-1240, 2001
                                                                                                                                                                                        RESULT 37
A89761
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hypothetical protein C05C10.4 - C
C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_rev
C;Accession: T18945
R;Matthews, P.
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T18945
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A;Map position: 2
A;Introns: 31/3; 96/3; 128/1; 155/3; 185/3;
A;Introns: 31/3; 96/3; 128/1; 155/3; 185/3;
C;Superfamily: mammalian acid phosphatase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Experimental C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              submitted to the EMBL Data Library, February 1995 A; Reference number: 219049
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    片
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A;Introns: 31/1; 51/2; 60/3; 76/2; 106/3; 163/1; 191/2; 226/3; 252/1; 274/3; 298/1; A;Note: T25B15.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Alcaraz, J.P.; Clabault, submitted to the Protein S A;Reference number: Z23021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hypothetical protein T25B15.30 - Arabidopsis thaliana (;Species: Arabidopsis thaliana (mouse-ear cress) C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 04-Feb-2000 #sequence_revision 04-Feb-2000 #text_change 09-Jul-2004 C;Accession: T46096
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: UNIPROT:Q09451; A;Experimental source: clone C05C10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Accession: T18945
A;Status: preliminary; translated from
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A;Experimental source: cultivar Columbia; BAC clo
  A;Title: Whole
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Residues:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: DNA
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Best Local
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5; Conserva
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5; Conserv
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genome sequencing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88
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83.3%;
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Sequence Database, January 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         62.5%;
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of meticillin-resistant Stapylococcus aureus
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Pred. No.
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Pred. No.
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le+02;
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K.; F
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A;Molecule type: DNA
A;Residues: 1-431 <KUR>
A;Cross-references: UNIPROT:Q9XB68; GB:BA000018; PID:g13699942; PIDN:BAB41241.1; GSPDB:A;Experimental source: strain N315
 밁
                                                                                                                                                                                                   A;Reference number: Z18831
A;Accession: T18221
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-492 <BAR>
                                                                                                                                                                                                                                                                                      R; Barrell, B.G.; Rajandream, M.A. submitted to the EMBL Data Library, November 1998
                                                                                                                                                                                                                                                                                                                      chromosome condensation regulator protein - yeast (Candida albicans)
C;Species: Candida albicans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 09-Jul-2004
C;Accession: T18221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Molecule type: DNA
A;Residues: 1-440 <1TO>
A;Cross-references: UNIPROT:Q9XB68; EMBL:D86934; PIDN:BAA82240.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Ito, T.; Katayama, Y.; Hiramatsu, K.
Antimicrob. Agents Chemother. 43, 1449-1458, 1999
A;Title: Cloning and nucleotide sequence determination of A;Reference number: Z22733; MUID:99278010; PMID:10348769
A;Accession: T44138
A;Status: preliminary; translated from GB/EMBL/DDBJ
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A;Accession: A89761
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Matches 5
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 HSYSKLP 37
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                               HOYSKLP 7
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1; Mismatches
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Pred. No.
                                                               Score 35; DB 2;
Pred. No. 1.3e+02;
0; Mismatches 1
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RESULT 40

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cytokine-inducible SH2 protein 4 - human
C;Species: Homo sapiens (man)
C;Date: 24-Uan-1998 #sequence_revision 13-Mar-1998 #text_change 09-Jul-2004
C;Accession: JC5762
R;Masuhara, M.; Sakamoto, H.; Matsumoto, A.; Suzuki, R.; Yasukawa, H.; MitsuBiochem. Biophys. Res. Commun. 239, 439-446, 1997
A;Title: Cloning and characterization of novel CIS family genes.
A;Reference number: JC5760; MUID:98008857; PMID:9344848
A;Accession: JC5762
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                                                                                                                                                               A;Cross-references: UNIPROT:O14544; DDBJ:AB006968; NID:g2463524; PIDN:BAA22538.1; C;Comment: This protein plays a role in the negative regulation of cytokine signal F;384-461/Domain: SH2 homology <SH2>
                                                                                                                                                                                                                                 A; Molecule type: mRNA
A; Residues: 1-535 < MAS>
                                                                                                                                                                                                                                                                          A;Status: nucleic acid sequence not shown
                                                                                   Matches
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Best Local (
  130 HHYSPAPW 137
                                         1 HQYSKLPW 8
                                                                                Similarity 5; Conserv
                                                                                Conservative
                                                                                                      62.5%;
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                                                                                   Score 35; DB Pred. No. 1.4e 0; Mismatches
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Search completed: December 17, 2004, 18:30:16
Job time : 3.53933 secs

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Result
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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1: uniprot_sprot:*
2: uniprot_trembl:*
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Gapop 10.0 , Gapext
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192.513 Million cell updates/sec
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AAS51605
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CHLE HUMAN
CHLE MOUSE
O90ZK8
BAC34196
O6MI24
CAE78158
O8CU788
MCE REOVD
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AAR11062
AAR11061
AAR11071
KV1B_HUMAN
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Compugen Ltd.
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Aas51605 ashbya go
P21927 oryctolagus
O9jkc1 rattus norv
P06276 homo sapien
O03311 mus musculu
Q90zk8 gallus gall
Bac34196 mus muscu
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P01610 homo sapien
QBysa0 anabaena sp
Q9vst8 drosophila
Q9x8f0 streptomyce
Q7naa3 photorhabdu
Q9gkj6 sus scrofa
Q13629 schizosacch
Q6fgg0 candida gla
Q89ug7 bradyrhizob
Q75b5 ashbya goss
                                           Bac34196 mus muscu
Q6mi24 bdellovibri
Cae78158 bdellovib
Q8my38 papilio xut
Q6cjy8 kluyveromyc
P11079 reovirus ty
Q91ra1 reovirus ty
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P01594 homo sapien
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Q9t0y1 lactobacill
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   homo sapien
RESULT Q9T0Y1 ID Q9 DT Q9 DT 011 DT 01 DT 01 DT 01 DT 01 DT PH PU COX LABOOC VILATION NC NC [1]
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37	37	37	37	37	37	37	37	37	37	37	37	37	37
66.1	66.1	66.1	66.1	66.1	66.1	66.1	66.1	66.1	66.1	66.1	66.1	66.1	66.1
585	566	444	402	402	388	330	330	245	242	190	134	134	108
N	N	N	N	۳	N	N	N	N	N	N	Ŋ	۲	N
Q9M2S5	Q7QFL1	056955	Q46434	OXA1_YEAST	Q9FLA1	CAE78520	Q6MQC7	Qauur6	Q9PU09	085999	Q8VDD0	KV4C_HUMAN	Q9CTV5
Q9m255 arabidopsis	07qfl1 anopheles g	O56955 human papil	Q46434 chlamydia t	P39952 saccharomyc	Q9fla1 arabidopsis	Cae78520 bdellovib	Q6mqc7 bdellovibri	Q8uut6 brachydanio	Q9pu09 latimeria c	O85999 sphingomona	_	P06314 homo sapien	Q9ctv5 mue mueculu

## ALIGNMENTS

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Q9JL84;
01-OCT-2000
01-OCT-2000
01-OCT-2003
O9TOY1 PRELIMINARY; PRT; 770 AA.

O9TOY1;
O1-MAY-2000 (TrEMBLrel. 13, Created)
O1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
O1-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Putative primase.
Putative primase.
Lactobacillus casei bacteriophage A2.
Viruses; dsDNA viruses, no RNA stage; Caudovirales; Siphoviridae.
NCBI TaxID=51369;
[1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPRO07110; Ig-like.
InterPro; IPRO03596; Ig v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
SMCNTE; PS50835; IG_LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Malkiel S., Liao L., Cunningham M.W., Diamond B.;
"T-Cell-dependent antibody response to the dominant epitope of
streptococcal polymaccharide, N-acetyl-glucomamine, is cross-reactive
with cardiac myosin.";
with cardiac myosin.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus (Mouse)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Infect. Immun. 68:5803-5808(2000).
EMBL; AF206022; AAF69320.1; -.
HSSP; P01594; 1JV5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. STRAIN=BALB/c;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=10090;
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(Fragment).
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                                                                                                                                                                                                                                                                                                   2 QYSKLPWT 9
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                                                                                                                                                                                                                                                                             QYSKFPWT 97
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107 AA;
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(TrEMBLrel. 15, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
immunoglobulin light chain variable region
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Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                    11648 MW; ACF9B1253ACA1E5D CRC64;
                                                                                                                                                                                                                                                                                                                                                                               78.6%;
87.5%;
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                                                                                                                                                                                                                                                                                                                                                       ; Score 44; DB 2
; Pred. No. 3.6;
0; Mismatches
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Matches 5
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                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
Ladero V., Garcia P., Alonso J.C., Suarez J.E.;
"A2 Cro, the lysogenic cycle repressor, specifically binds
genetic switch region of Lactobacillus casei bacteriophage
Virology 262:222-229(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.

MEDLINE=97197195; PubMed=9044284;

MEDLINE=97197195; PubMed=9044284;

Garcia P., Alonso J.C., Suarez J.E.;

"Molecular Analysis of the cos Region of the Lactobacillus casei

Bacteriophage A2. Gene product 3, gp3, specifically binds to its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bacteriophage A2.";
J. Bacteriol. 180:3474-3476(1998).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A. MEDLINE=98317291;
                                                                                                                                                                                                                                 SEQUENCE FROM N.A.

Garcia P., Ladero V., Suarez J.E.;

Submitted (MAY-2002) to the EMBL/GenBank/DDBJ

EMBL; AJ251789; CAB63672.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Garcia P., Ladero V., Alonso J.C., Suarez J.E.; "Cooperative interaction of CI protein regulates lysogeny of Lactobacillus casei by bacteriophage A2."; J. Virol. 73:3920-3929(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Moscoso M., Suarez J.E.;
"Characterisation of the DNA replication module of bacteriophage A2
"Characterisation of replication as a defence against infection
and use of its origin of replication as a defence against infection
during milk fermentation by Lactobacillus casei.";
Virology 273:101-111(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIINE=98445458; PubMed=9770432;
Alvarez M.A., Herrero M., Suarez J.E.;
"The Site-Specific Recombination system of the Lactobacillus spp
Bacteriophage A2 Integrates in Gram Positive and Gram Negative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                            TIGRFAMs;
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MEDLINE=99214330;
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                                                                                                                                                                                                     InterPro; IPR004968; Pox_D5.
InterPro; IPR006500; Primase_C.
                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=22262118; PubMed=12374837;
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                                                                                                                                                        PF03288; Pox D5; 1.
AMs; TIGR01613; primase_Cterm;
NCE 770 AA; 88233 MW; 382B
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                                                                                                     Similarity 5; Conser
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                                                                     HOYSKLPWT 9
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piol. 23:505-514(1997).
                                                                                                   76.8%;
larity 55.6%;
Conservative
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P., Bascaran V., Herrero M., Alvarez M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PubMed=10196287;
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                                                                                        Pred. No. ....
4; Mismatches
                                                                                                   4;
                                                                                                                               Score 43;
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Lafontaine I., de Montigny J., Marck C., Newveglise C., Talla E.,
A Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
A Barnay S., Blanchin S., Beckerich J.M., Beyne E., Bleykasten C.,
A Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
Bespons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
A Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
A Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
A Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
A Nicaud J.M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
Potler S., Richard G.F., Straub M.L., Suleau A.,
Pellenz S., Potler S., Richard G.F., Straub M.L., Suleau A.,
Pellenz S., Potler S., Richard G.F., Straub M.L., Suleau A.,
Pellenz C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
A Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
A Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
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Query Match
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AAR11015;
02-MAR-2004 (TrEMBLrel. 2
02-MAR-2004 (TrEMBLrel. 2
02-MAR-2004 (TrEMBLrel. 2
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ORFNames-DEHAOF0145g;
Debaryomyces hansenii (Yeast) (Torulaspora hansenii).
Bebaryomyces hansenii (Yeast) (Torulaspora hansenii).
Bukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Debaryomyces.
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NON TER
SEQUENCE
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01-OCT-2004 (TrEMBLrel. 28,
01-OCT-2004 (TrEMBLrel. 28,
01-OCT-2004 (TrEMBLrel. 28,
Similarities with sp|P47168
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                                                                            Genoscope;
Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases
EMBL; CR382138; CAG88702.1; -.
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Mammalia; Eutheria; Rodentia;
                                                      SEQUENCE
                                                                                                                                                           STRAIN=CBS767;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENOLEVURES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=CBS767;
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Liang Z., Xie C., Chen C., Kre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=10090;
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                                                                                                                                                                                       SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                 Genome evolution
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MAR-2004 (TrEMBLrel. 27, Last annotation update)
immunoglobulin kappa light chain (Fragment).
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107 AA;
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                                                      A
                                                                                                                                                                                                                                                                 in yeasts.";
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                                                                                                                                                                                                                                                                                               J.L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11408 MW;
                                                      65217 MW;
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Sciurognathi;
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                                                         73D939503DED2E8D CRC64;
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No.
     41;
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13;
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thi; Muridae; Murinae; Mus
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KV1R_HUMA
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GO; GO:0008415; F:acyltransferase activity; IE
GO; GO:000840; F:gamma-glutamyltransferase ac
GO; GO:0003840; F:transferase activity; IEA.
InterPro; IPR000101; GGT_peptidase.
Pfam; PF01019; Ggluttranspept; 1.
Pfam; PF01019; Ggluttranspept; 1.
PRINTS; PR01210; GGTRANSPTASE.
ACYltransferase; Complete proteome; Transferase; Acyltransferase; Complete proteome; Transferase; SEQUENCE 623 AA; 65009 MW; 2201015F85FEA5:
                                                                                                                                                                                                                                                                                                                                                                                          KV1R HUMAN
P01610;
21-JUL-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Salanoubat M., Génin S., Artiguenave F., Gouzy J., Mangenot S., Arlat M., Billault A., Brottier P., Camus J.C., Cattolico L., Chandler M., Choisne N., Claudel-Renard C., Cunnac S., Demange N., Gaspin C., Lavie M., Woisan A., Robert C., Saurin W., Schiex T., Siguier P., Thebault P., Whalen M., Wincker P., Levy M., Weissenbach J., Boucher C.A.; "Genome sequence of the plant pathogen Ralstonia solanacearum."; Nature 415:497-502(2002).

EMBL, AL646072; CAD16628.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Q8XVB0
Q8XVB0;
01-MAR-2002
01-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HUMAN
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01-MAR-2002 (TrEMBLrel. 20, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
PROBABLE GAMMA-GLUTAMYLTRANSPEPTIDASE PROTEIN (EC 2
                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                       21-JUL-1986 (Rel.
21-JUL-1986 (Rel.
05-JUL-2004 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Name=ggt2; Synonyms=RS00172; OrderedLocusNames=RSc2921;
Ralstonia solanacearum (Pseudomonas solanacearum).
Bactoria; Proteobacteria; Betaproteobacteria; Burkholderiales;
MEDLINE-83273707; PubMed=6410398; Goni F., Frangione B.; Goni F., Frangione B.; "Amino acid sequence of the Fv region of a human monoclonal I (protein WEA) with antibody activity against 3,4-pyruvylated in Klabsiella polysaccharides K30 and K33."; Proc. Natl. Acad. Sci. U.S.A. 80:4837-4841(1983).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Burkholderiaceae; Ralstonia.
                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                 NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                       Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=21681879; PubMed=11823852;
                                                                                                                                                                                                                                                                                                               Ig kappa chain V-I region WEA.
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01, Last sequence update)
44, Last annotation updat
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l; Mismatches
                                                                                                                                                                                                                                    Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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2201015F85FEA576
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                                                                                                                      EMBL; AP003592; BAB74888.1; -.

PIR; AF2204; AF2204.

GO; GO:0016701; F:oxidoreductase activity, ac

GO; GO:0016119; F:carotene metabolism; IEA.

InterPro; IPR011393; Carotene_ketolas.

InterPro; IPR05804; Fa_desat.

InterPro; IPR010257; FA_desat_sub.

Pfam; PF00487; FA_desat_trase; 1.

PIRSF; PIRSF027840; Carotene_ketolas; 1.

Probom; PD001081; FA_desat_sub; 1.
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SEQUENCE FROM N.A.

MEDLINE=21595285; PubMed=11759840;

Kaneko T., Nakamura Y., Wolk C.P., Kuritz T., Sasamoto S.,

Kaneko T., Nakamura Y., Wolk C.P., Kuritz T., Sasamoto S.,

Kaneko T., Nakamura Y., Wolk C.P., Kuritz T., Sasamoto S.,

Matanabe A., Iriguchi M., Ishikawa A., Kawashima K., Kimura

Matanabe A., Nuraki A.,

Matanabe A., Muraki A.,

Matanabe A., Yamada M.,
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SEQUENCE
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01-MAR-2002 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
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InterPro; IPR003596; Ig v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
SPROSITE; PS50835; IG_LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                  "Complete genomic sequence of the filamentous nitrogen-fixing cyanobacterium Anabaena sp. strain PCC 7120.";
DNA Res. 8:205-213(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Watanabe A., Iriguchi M., Ishikawa A.
Kishida Y., Kohara M., Matsumoto M.,
Nakazaki N., Shimpo S., Sugimoto M.,
Yasuda M., Tabata S.;
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HSSP, P80352; IWTL.
GO: GO:0005576; C:extracellular; NAS.
GO; GO:0003823; F:antigen binding; NA
GO; GO:0006955; P:immune response; NA
                                                                                                      Complete proteome
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Bacteria; Cyanobacteria; Nostocales;
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Q9VST8;
01-MAY-2000
01-MAY-2000
           MEDINE-24.26065; PubMed=12537568;

Celniker S.E., Wheeler D.A., Kronmiller B., Carlson J.W., Halpe Patel S., Adams M., Champe M., Dugan S.P., Frise E., Hodgson A. George R.A., Hoskins R.A., Laverty T., Muzny D.M., Nelson C.R., Pacleb J.M., Park S., Pfeiffer B.D., Richards S., Sodergren E.J. Pacleb J.M., Park S., Wan K., Stapleton M., Sutton G.G., Ver Weinstock G., Scherer S.E., Wyers E.W., Gibbs R.A., Rubin G.M., "Finishing a whole-genome shotgun: release 3 of the Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORFNames=CG4942;
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           Q9X8F0;
01-NOV-1999 (TrEMBLrel. 12, C:
01-NOV-1999 (TrEMBLrel. 24, L:
01-JUN-2003 (TrEMBLrel. 24, L:
Putative phosphotransferase.
OrderedLocusNames=SC03277; OR:
Streptomyces coelicolor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Stapleton M., Brokstein P., Hong L., Agpayani A., Champe M., Chavez C., Dorsett V., Farfan D., Frise E., Glonzalez M., Guarin H., Li P., Liao G., Miranda A., Mung. Nunco J., Pacleb J., Paragas V., Park S., Phouanenavong Yu C., Lewis S.E., Rubin G.M., Celniker S.; Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misra S., Crosby M.A., Mungall C.J., Matthews B.B., Campbell K.S., Hradecky P., Huang Y., Kaminker J.S., Millburn G.H., Prochnik S.E. Smith C.D., Tupy J.L., Whitfied E.J., Bayraktaroglu L., Berman B.P Bettencourt B.R., Celniker S.E., de Grey A.D., Drysdale R.A., Harris N.L., Richter J., Russo S., Schroeder A.J., Shu S.O., Stapleton M., Yamada C., Ashburner M., Gelbart W.M., Rubin G.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- SUBCELLULAR LOCATION: Integral membrane protein
-!- SIMILARITY: Belongs to the OXA1/oxaA family.
EMBL; AE003553; AAF50323.1; -.
EMBL; AY069618; AAL39763.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patel S., Frise E., Wheeler D.A., Lewis S.E., Rubin G.M., Ashburner M., Celniker S.E.; "The transposable elements of the Drosophila melanogaster euchromatin: a genomics perspective."; a genome Biol. 3:RESEARCH0084-RESEARCH0084(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pfam; PF02096; 60KD_IMP; 1.
Transmembrane
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FlyBase; FBgn00
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SEQUENCE 351
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GO:0007165; P:signal transduction;
erPro; IPR001708; 60kDa_innermeb.
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O1-MAR-2004 (TrEMBLrel. 26, Created)
O1-MAR-2004 (TrEMBLrel. 26, Last sequence update)
O1-MAR-2004 (TrEMBLrel. 26, Last annotation updat
Complete genome; segment 1/17.
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Thomson N.R., James K.D., Harris D.E., Quail M.A., Kieser H.,
Harper D., Bateman A., Brown S., Chandra G., Chen C.W., Collins M.,
Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth S.,
Huang C.-H., Kieser T., Larke L., Murphy L.D., Oliver K., O'Neil S.,
Rabbinowitsch E., Rajandream M.A., Rutherford K.M., Rutter S.,
Seeger K., Saunders D., Sharp S., Squares R., Squares S., Taylor K.,
Warren T., Wietzorrek A., Woodward J.R., Barrell B.G., Parkhill J.,
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Nature 417:141-147(2002).
EMBL; AL939115; CAB40335.1; -.
PIR; T36237; T36237.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=22957627; PubMed=14528314;
Duchaud E., Rusniok C., Frangeul L., Buchrieser C., Givaudan A.,
Taourit S., Bocs S., Boursaux-Ende C., Chandler M., Charles J.-F.,
Dassa E., Derose R., Derzelle S., Freyssinet G., Gaudriault S.,
Medigue C., Lanois A., Powell K., Siguier P., Vincent R., Wingate V.,
Zouine M., Glaser P., Boemare N., Danchin A., Kunst F.;
"The genome sequence of the entomopathogenic bacterium Photorhabdus
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Complete proteome; T
SEQUENCE 288 AA;
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PhotoList, plu0031; -.
InterPro; IRR000276; GPCR Rhodpsn.
PROSITE; PS00237; G_PROTEIN_RECEP_F1_1; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=TT01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Photorhabdus luminescens (subsp. laumondii).
Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OrderedLocusNames=plu0031;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q7NAA3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Streptomycineae;
NCBI_TaxID=1902;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=141679;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enterobacteriaceae; Photorhabdus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro;
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InterPro; IPR002575; APH_trans.
InterPro; IPR011009; Kinase_like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hopwood D.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             luminescens.";
                                                                                                                                                                                                                                                                                                                                                                                                                    Biotechnol. 21:1307-1313(2003).; BX571859; CAE12326.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    256
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QYSKLPWT
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                                                                                                                                                                                                                                  38412 MW;
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75.0%;
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75.0%;
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                                                                                                         Score 38;
Pred. No.
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Pred. No. 1.
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                                                                                                                                                                                                                              7EB8746696E2B8B6 CRC64;
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                                                                                                         1.4e+02;
                                                                                                                                                     DB 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        update)
                                                                                                                                                 Length 336;
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Best Local (
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01-JAN-1998
01-JAN-1998
05-JUL-2004
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01-MAR-2001
01-MAR-2001
01-OCT-2003
                                                                                                                                                                                                                                                                                                                               01-JAN-1998 (TrEMBLrel. 05, Created)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
LIPOATE-PROTEIN LIGASE A (SPBC17A3.09c protein) (Pi037 protein)
Name=pi037; Synonyms=SPBC17A3.09c;
Schizosaccharomyces pombe (Fission yeast).
Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yanagida M.;
"A 38 kb segment containing the cdc2 gene from the left arm yeast chromosome II: sequence analysis and characterization genomic DNA and cDNAs encoded on the segment.";
                                                                                                          MEDLINE=20089027; PubMed=10620777;
Machida M., Yamazaki S., Kunihiro S.,
Haikawa Y., Yamazaki J., Yamamoto S.,
Sakai M., Aoki K., Ogura K., Kudoh Y.,
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InterPro; IPR000997; Cholinesterase.
InterPro; IPR000379; Ser estrs.
Pfam; PF00135; COesterase; 1.
PRINTS; PR00878; CHOLNESTRASE.
PROSITE; PS00122; CARBOXYLESTERASE_B_1;
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MEDLINE=21419000; PubMed=11528129;
Verle M., Tuggle C.,
  genomic DNA and cDNAs
Yeast 16:71-80(2000).
                                                                                                                                                                                                    SEQUENCE FROM N.A.
STRAIN=972 h-;
                                                                                                                                                                                                                                                            NCBI_TaxID=4896;
                                                                                                                                                                                                                                                                                            Schizosaccharomyces.
                                                                                                                                                                                                                                                                                                                 Schizosaccharomycetales;
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Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.
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6; Conserv
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(TrEMBLrel. 16, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
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75.0%;
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Pred. No.
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                                                                                                          , Tanaka T., Ku
, Sekine M., Og
., Kikuchi H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      363
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                                                                                                          , Kushida N.,
, Oguchi A., N
H., Zhang M.Q.
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RESULT 13
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RA Sgouros J., Peat N., Hayles J., Baker S., Basham D., Bowman S.,
RA Brooks K., Brown D., Brown S., Chillingworth T., Churcher C.,
RCOLlins M., Connor R., Cronin A., Davis P., Feltwell T., Fraser A.,
RA Gollins M., Connor R., Cronin A., Davis P., Feltwell T., Fraser A.,
RA Holroyd S., Hornsby T., Howarth S., Huckle E.J., Hunt S., Jagels K.,
RA Holroyd S., Hornsby T., Howarth S., Huckle E.J., Hunt S., Jagels K.,
RA Holroyd S., Hornsby T., Howarth S., Huckle E.J., Hunt S., Jagels K.,
RA Holroyd S., Hornsby T., Howarth S., Huckle E.J., Hunt S., Jagels K.,
RA Holroyd S., Hornsby T., Howarth S., Huckle E.J., Hunt S., Jagels K.,
RA Mooney P., Moule S., Mangall K., Murphy L., Niblett D., Odell C.,
RA Mooney P., Moule S., Mangall K., Murphy L., Niblett D., Odell C.,
RA Rutherford K., Rutter S., Saunders D., Seeger K., Sharp S.,
RA Rutherford K., Rutter S., Saunders D., Seeger K., Sharp S.,
RA Waldien J., Volckeert G., Aert R., Roben J., Grymonprez B.,
RA Waldiens I., Vanstreels E., Rieger M., Schafer M., Muller-Auer S.,
RA Weltjens I., Vanstreels E., Rieger M., Schafer M., Moestl D.,
RA Hilbert H., Borzym K., Langer I., Beck A., Lehrach H., Reinhardt R.,
RA Hilbert H., Borzym K., Langer I., Beck A., Lehrach H., Reinhardt R.,
RA Hilbert F., Aves S.J., Xiang Z., Hunt C., Moore K., Hurst S.M.,
RA Galibert F., Aves S.J., Xiang Z., Hunt C., Moore K., Hurst S.M.,
RA Galibert F., Aves S.J., Jimenez J., Sanchez M., del Rey F., Benito J.,
RA Lucas M., Rochet M., Gaillardin C., Tallada V.A., Garzon A., Thode G.,
RA Daga R.R., Cruzado L., Jimenez J., Sanchez M., del Rey F., Benito J.,
RA Crutti L., Lowe T., McCombie W.R., Paulsen I., Potashkin J.,
The genome sequence of Schizosaccharomyces pombe.";
RMIL, Abno1457. naa514171.
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Best Local S
Matches 5
                                                                                                                                                                                                                                                                      Q6FQG0 PRELIMINARY; PRT; 386 AA. Q6FQG0; Q6FQG0; Q5-JUL-2004 (TrEMBLrel. 27, Created) 05-JUL-2004 (TrEMBLrel. 27, Last sequence update) 05-JUL-2004 (TrEMBLrel. 27, Last annotation update) 5-JUL-2004 (TrEMBLrel. 27, Last sequence.
                                                                                                                          Candida glabrata (Yeāst) (Torulopsis glabrata).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; mitosporic Saccharomycetales; Candida.
SEQUENCE FROM N.A. STRAIN=CBS138;
                                                                                            Eukaryota; Fungi; Ascomycota; Saccharomycetales; mitosporic NCBI_TaxID=5478;
                                                                                                                                                                                                                                                ORFNames=CAGL0I06534g;
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MEDLINE=21848401;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=972
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SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GO:0016874; F:ligase activity; IEA.
GO:0016874; F:ligase activity; IEA.
GO:0006464; P:protein modification; IEA.
EarPro; IPR004143; BPL LipA LipB.
EARPAMB; TIGR00545; lippyltrans;
RFAMB; TIGR00545; lippyltrans; 1.
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55.6%;
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Pred. No. 1.5e+02;
2; Mismatches 2
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Matches 6
                                                                                                                                              Bradyrhizobium japonicum USDA110.";
DNA Res. 9:189-197(2002).
EMBL; AD005940; BAC46715-1; -.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0005215; F:transport; IEA.
GO; GO:0006510; P:transport; IEA.
InterPro; IPR001851; Bac inmem_transp.
Pfam; PF02653; BPD transp_2; 1.
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Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla E.,
Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
Barnay S., Blanchin S., Beckerich J.M., Beyne E., Bleykasten C.,
Bairame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
Begisrame A., Boyer J., Cattolico L., Ferry-Dumazet H., Groppi A.,
Boisrame A., Hone E., Fairhead C., Ferry-Dumazet H., Groppi A.,
Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
Kerrest A., Koszul R., Lemaire M., Lesur I., Ma L., Muller H.,
Nicaud J.M., Nikolski M., Oztas S., Ozier-Kalogeropoulos O.,
Pellenz S., Potier S., Richard G.F., Straub M.L., Suleau A.,
Swennene D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
Zeniou-Meyer M., Zivanovic I., Bolotin-Fukuhara M., Thierry A.,
Bouchier C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
Winneyer B., Cariet J. T.,
Winneyer B., Scarpelli C., Gaillardin C., Weissenbach J.,
                                                                                                                     Complete
SEQUENCE
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-!- SIMILARITY: Belongs to the OXA1/oxaA family.
EMBL; CR380955; CAG60471.1; -.
InterPro; IPR001708; 60kDa innermeb.
Pfam; PF02096; 60KD_IMP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bradyrhizobium japonicum.
Bacteria; Proteobacteria; Alphaproteobacteria; Rhizobiales;
Bradyrhizobiaceae; Bradyrhizobium.
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01-JUN-2003 (TrEMBLrel. 24,
01-OCT-2003 (TrEMBLrel. 25,
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Dujon B., Sh
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MEDLINE=22484998; PubMed=12597275;
Kaneko T., Nakamura Y., Sato S., Minamisawa K., Uchiumi T.,
Sasamoto S., Watanabe A., Idesawa K., Iriguchi M., Kawashima K.
Kohara M., Matsumoto M., Shimpo S., Tsuruoka H., Wada T., Yamad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OrderedLocusNames=blr1450;
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01-JUN-2003
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Bouchier P., Souciet J.L.;
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                                                                                                                                                                                                                                                                                       'Complete genomic sequence of nitrogen-fixing symbiotic bacterium
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                                                    Conservative
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75.0%;
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Pred. No. 1.6e+02
                                                                                                                                                                                                        activity;
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                                                                                                                     2C7D90C36B8E1B26 CRC64;
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RESULT 16
AAS5165
ID AAS51
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CT 23-AP
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05-JUL-2004
05-JUL-2004
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Q75B85;
                                                                                                                                                                                                                                                                                                                   PROSITE; PS00107; PROTEIN_KINASE_ATP; 1.
PROSITE; PS50011; PROTEIN_KINASE_DOM; 1.
PROSITE; PS00108; PROTEIN_KINASE_ST; UNKNOWN_1.
ATP-binding; Kinase; Transferase.
SEQUENCE 434 AA; 49672 MW; E3F62F511479CF1B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL;
                                                                                                                                                                                                                                                                                                                                                                                   ProDom; PD000001; Prot_kinase; 1.
SMART; SM00220; S_TKC; 1.
SMART; SM00219; TyrKC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPRO11009; Kinase like.
InterPro; IPRO00719; Prot kinase.
InterPro; IPR002290; Ser_thr_pkinase.
InterPro; IPR008271; Ser_thr_pkin_AS.
InterPro; IPR001245; Tyr_pkinase.
Pfam; PP00069; Pkinase; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ashbya gossypii (Yeast) (Eremothecium gossypii).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Eremothecium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (DEC-2002) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A. STRAIN=ATCC 10895;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PubMed=15001715;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN-ATCC 10895;
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 SEQUENCE FROM N.A.
                                   Saccharomycetales;
                                                 Ashbya gossypii (Yeast) (Eremothecium gossypii).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
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                     NCBI_TaxID=33169;
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23-APR-2004
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                                                                                                                                                                                                                                                                     Similarity 5; Conser
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                                                                                                                                                                                                                                           QYSKLPW 8
                                              Fungi; Ascomycota;
                                                                                                  (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
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                                     Saccharomycetaceae;
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27,
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Last annotation update)
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2; Mismatches
                                                                                                                          Created)
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                                      Eremothecium.
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PubMed=15001715;
Dietrich F.S., Voegeli S., Brachat S., Lerch A., Gates K., Ste Dietrich F.S., Voegeli S., Brachat S., Lerch A., Gates K., Ste Mohr C., Pohlmann R., Luedi P., Choi S., Wing R.A., Flavier A. Gaffney T.D., Philippen P.;
"The Ashbya gossypii genome as a tool for mapping the ancient Saccharomyces cerevisiae genome.";
Science 304:304-307(2004).
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01-MAY-1991 (Rel. 18, Last sequence up
01-OCT-2004 (Rel. 45, Last annotation
Cholinesterase precursor (EC 3.1.1.8)
(Choline esterase II) (Butyrylcholine
                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
STRALN=New Zealand;
MEDLINE=90326526; PubMed=2374720;
Jbilo O., Roudani S., Chatonnet A.;
"Complete sequence of rabbit butyrylcholinesterase.";
Nucleic Acids Res. 18:3990-3990(1990).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P21927;
01-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN=ATCC 10895;
                                                                                                                                                                                                                                                                                                                                                       MEDLINE=91201348; PubMed=2016308;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Name=BCHE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Philippsen P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=ATCC 10895;
                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE OF 75-215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pseudocholinesterase)
                                                                                                                                two subunits in a dimer are linked by a disulfide bond. TISSUE SPECIFICITY: Present in most cells except erythrocytes. MISCELLANEOUS: Cholinesterase is highly reactive with organophosphate esters.
SIMILARITY: Belongs to the type-B carboxylesterase/lipase family.
                                                                                                                                                                                                                     carboxylic acid anion.
SUBUNIT: Homotetramer. The tetramer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RABIT
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5; Conserve
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71.4%;
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                                                                                                                                                                                                                                                                                                                                                                                        N.A.
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Pred. No.
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(Acylcholine acylhydrolase)
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EMBL; X52091; CAA36308.1;
EMBL; X52092; CAA36308.1;
EMBL; M62779; AAA31169.1;
                       HSSP;
GO; GO
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Brimijoin S., Hinrichs S
Submitted (MAR-2000) to t
                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
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PROSITE; PS00122; CARBOXYLESTERASE_B_
PROSITE; PS00941; CARBOXYLESTERASE_B_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Butyrylcholinesterase
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                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                  TISSUE=Heart;
                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rattus norvegicus
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GO:0003824; F:catalytic activity; GO:0004104; F:cholinesterase activerPro; IPR002018; CarbesteraseB.
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Rodentia;
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75.0%;
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Charge relay system (By s:
By similarity.

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1; Mismatches
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Lockridge O.;
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Cholinesterase.
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Sciurognathi; Muridae; Murinae; Rattus.
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                         activity;
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Best Local S
Matches 6
                                                                                                                                                                                                                           Kott M., No.
MEDLINE-2388257; PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Strausberg R.D., Collins F.S., Wagner L. Shemmen C.M., Schuler G.D., Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.Fr., Bhat N.K., Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Villalon B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-OCT-2004 (Rel. 45, Last annotation Cholinesterase precursor (EC 3.1.1.8) (Choline esterase II) (Butyrylcholine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROSITE; PS00122; CARBOXYLESTERASE_B_1; PROSITE; PS00941; CARBOXYLESTERASE_B_2; Hydrolase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR000997; Cholinesterase.
InterPro; IPR000379; Ser estrs.
Pfam; PF00135; COesterase; 1.
PRINTS; PR00878; CHOLNESTRASE.
                                                                                                                                                                                                                                                                                                                            MEDLINE=87231856; PubMed=3035536;
Prody C.A., Zevin-Sonkin D., Gnatt A., Goldberg O
"Isolation and characterization of full-length cD
cholinesterase from fetal human tissues.";
Proc. Natl. Acad. Sci. U.S.A. 84:3555-3559(1987).
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01-AUG-1988
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                                                                                                                                                                             SEQUENCE FROM
TISSUE=Skin;
                                                                                                                                                                                                                                                           McTiernan
                                                                                                                                                                                                                                                                     MEDLINE=88016155; PubMed=3477799;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Biochemistry
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (Human)
Eukaryota, Metazoa, (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Name=BCHE; Synonyms=CHE1;
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                                                                                                                                                                                                                  n C., Adkins S., Chatonnet A., Vaughan T.A., Rosenberry T.L., la Du B.N., Lockridge O.; DNA clone for human cholinesterase."; tl. Acad. Sci. U.S.A. 84:6682-6686(1987).
                                                                                                                                                                                                                                                                                                                                                                                                            FROM
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8 (Rel. 08, Last &
4 (Rel. 45, Last &
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75.0%;
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Pred. No.
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EMBL; M32391; J
EMBL; M32389; J
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EMBL; M16541; J
EMBL; M16474; J
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Moriwaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D. Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.; "Generation and initial analysis of more than 15,000 full-l and mouse CDNA sequences.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     McGuire M.C., Nogueira C.P., Bartels C.F., Lightstone H., F
van der Spek A.F.L., Lockridge O., la Du B.N.;
"Identification of the structural mutation responsible for
"dibudaine resistant (atypical) variant form of human serum
                                                                                                                                                            the
                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=96287386; PubMed=8680411; Iida S., Kinoshita M., Fujii H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Structure of human serum Bioessays 9:125-128(1988)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cholinesterase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=88007487; PubMed=3115973;
Lockridge O., Adkins S., la Du B.N.;
"Location of disulfide bonds within
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE-87109144; PubMed=3542989; Lockridge O., Bartels C.F., Vaugh
                                                                                               or send an
                                                                                                              entities requires a
                                                                                                                           modified and this statement is not removed.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              hypocholinesterasemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Mutations of human butyrylcholinesterase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cholinesterase."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lockridge O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=89149758; PubMed=3067729;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'Complete
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             omplete amino acid sequence of human serum cholinesterase.";
Biol. Chem. 262:549-557(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Biol.
                                                                                                                                                                                                                                                                                                                              carboxylic acid anion.
SUBUNIT: Homotetramer. The tetramer is composed of two dimers.
two subunits in a dimer are linked by a disulfide bond.
TISSUE SPECIFICITY: Present in most cells except erythrocytes.
DISEASE: Mutant alleles of CHE1 are responsible for
                                                                                                                                        European Bioinformatics Institute. The by non-profit institutions as long
                                                                                                                                                                                                                                                                                                                                                                                                             CATALYTIC ACTIVITY: An a
                                                                                                                                                                                                                     organophosphate esters.
SIMILARITY: Belongs to the type-B carboxylesterase/lipase family.
                                                                                                                                                                                                                                                                                hypocholinesterasemia resulting in suxamethonium sensitivity. Homozygous persons sustain prolonged apnea after administration the muscle relaxant suxamethonium in connection with surgical
                                                                                                                                                                                                                                                 anesthesia.
MISCELLANEOUS: Cholinesterase is highly reactive with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chem. 262:12945-12952(1987).
                                                                                            equires a license agreement (Semail to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Acad.
AAA99296.1; JOINED.
AAA99296.1; JOINED.
AAA99296.1; JOINED.
AAA98113.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   serum cholinesterase.";
                                                                                                                                                                                                                                                                                                                                                                                                               An acylcholine + H(2)O =
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vaughan T.A., Wong C.K., Norton S.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Moriyama Y., Nakamura Y., Yura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the
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                                                                                                                       There are no restrictions on ong as its content is in no oved. Usage by and for commen
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DE Cholinesteras
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Matches
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GO; GO:0001824; F:catalytic activity; NAS.
GO; GO:0004104; F:cholinesterase activity; NAS.
GO; GO:001989; F:enzyme binding; NAS.
GO; GO:0050783; P:cocaine metabolism; TAS.
InterPro; IPR002018; CarbesteraseB.
InterPro; IPR003019; Carbesterase.
InterPro; IPR00379; Ser_estrs.
Pfam; PF00135; COcosterase; I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PDB; iEHQ; Model; A=30-560.

PDB; 1P0I; X-ray; A=29-557.

PDB; 1P0M; X-ray; A=29-557.

PDB; 1P0P; X-ray; A=29-557.

PDB; 1P0Q; X-ray; A=29-557.
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CARBOHYD
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ACT_SITE
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ACT_SITE
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MIM; 177400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRINTS; PR00878; CHOLNESTRASE.

PROSITE; PS00122; CARBOXYLESTERASE_B_1; 1.

PROSITE; PS00941; CARBOXYLESTERASE_B_2; 1.

3D-structure; Direct protein sequencing; Disease
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; 1EHQ; Model;
; 1POI; X-ray;
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                                                                                                                                     Similarity
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         (Rel. 27,
(Rel. 27,
(Rel. 45,
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                                              STANDARD;
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N-linked (GlCNAc...).
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N-linked (GlCNAc...).
N-linked (GlCNAc...).
N-linked (GlCNAc...).
N-linked (GlCNAc...).
                                                                                                                                                                                                                                         D -> G (in atypical form, cresistant; dbsNp:1799807).
/FIId=VAR_002360.
T -> M (in fluoride-1)
                                                                                                                           Score 38; DB : Pred. No. 2.5e
                                                                                                                                                                         activity;
/FTId=VAR_
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Acyl-ester intermediate
Acyl-ester intermediate
Charge relay system (By
Charge relay system (By
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G -> V (in fluoride-2)
/FTId=VAR_002363.
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L -> I (in hypocholinesterasemia)
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                                                                                                                                                                 C9836409D9057F27
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y similarity).
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Cholinesterase

precursor

Last sequence update)
Last annotation update)
rsor (EC 3.1.1.8) (Acylcholine acylhydrolase)

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Query Match
Best Local
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CARBOHYD
CARBOHYD
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CONFLICT
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MEDLINE=90380429; PubMed=2400605;

Rachinsky T.L., Camp S., Li Y., Ekstroem T.J., Newton M.,

"Molecular cloning of mouse acetylcholinesterase: tissue d

of alternatively spliced mRNA species.";

Neuron 5:317-327(1990).
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DISULFID
                                                                                                                                                                                                                                                                                                                                                                       Pfam; PrOU15; CHOLNESTRASE.

PRINTS; PRO0878; CHOLNESTRASE B 1; 1.

PROSITE; PS00122; CARBOXYLESTERASE B 2; 1.

PROSITE; PS00941; CARBOXYLESTERASE B 2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        J. Biol. Chem. 266:6966-6974(1991).
-!- CATALYTIC ACTIVITY: An acylcholine + H(2)O =
                                                                                                                                                                                                                                                                                                                                                             Glycoprotein; SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified and this statement is not removed entities requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no restues by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Arpagaus M., Chatonnet A., Masson P., Newton M., Vaughan T.A., Bartels C.F., Nogueira C.P., la Du B.N., Lockridge O.; "Use of the polymerase chain reaction for homology probing of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
                                                                                                                                                                                                                   DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; M99492; AAA37328.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    butyrylcholinesterase from several vertebrates.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=91201348; PubMed=2016308;
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SIMILARITY: Belongs to the type-B carboxylesterase/lipase family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MGI:894278; Bche.
rPro; IPR002018; CarbesteraseB.
rPro; IPR000997; Cholinesterase.
rPro; IPR000379; Ser_estrs.
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   Similarity
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                                                                                      Chordata;
Rodentia;
 67.9%;
75.0%;
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Interchain (By Similarity).

Interchain (GlCNAC...) (Potes N-linked (GlCNAC...) 
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Charge rela
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719B1B220D1E5367
 38;
No.
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relay system
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DB 1;
2.5e+02;
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HSSP; P21386; INSM.
GO; GO:0004104; F:cholinesterase activity; IE
GO; GO:0016787; F:hydrolase activity; IEA.
InterPro; IPR002018; CarbesteraseB.
InterPro; IPR000997; Cholinesterase.
InterPro; IPR000379; Ser_estrs.
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SEQUENCE FROM N.A.
STRAIN-C57BL/6J; TISSUE=Liver;
MEDLINE=235468; PubMed=12466851;
The FANTOM Consortium,
the RIKEN Genome Exploration Research Group Ph
"Analysis of the mouse transcriptome based on
60,770 full-length cDNAs.";
Nature 420:563-573(2002).
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01-DEC-2001 (TrEMBLrel. 19, Last sec
01-OCT-2003 (TrEMBLrel. 25, Last and
Butyrylcholinesterase precursor (EC
                                                                                                                                                                                                                                                                                      EUKaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=10090;
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Geissler K., Chatonnet A.,
Submitted (APR-2001) to th
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14-APR-2004 (TrEMBLrel. 27, Last sequence update)
14-APR-2004 (TrEMBLrel. 27, Last annotation update)
14-APR-2004 (TrEMBLrel. 27, Last annotation update)
Adult male liver tumor cDNA, RIKEN full-length enri
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PROSITE; PS00122; CARBOXYLESTERASE_B_1;
PROSITE; PS00941; CARBOXYLESTERASE_B_2;
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                                                                                                                                                                                                                                                                                                                                                                                                                              clone:C730038G20 product:butyrylcholinesterase.
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Veognathae; Galliformes; Phasianidae; Phasiani
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the EMBL/GenBank/DDBJ_databases.
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rsor (EC 3.1.1.8).
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RESULT 23
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C STRAIN-C57BL/6J; TISSUE-Liver;

K MEDLINE-20530913; PubMed=11076861;

X MEDLINE-20530913; PubMed=11076861;

X MEDLINE-20530913; PubMed=11076861;

X Shibata K., Itoh M., Alzawa K., Nagaoka S., Sasaki N., Carninci P.,

XA Shibata K., Itoh M., Alzawa K., Nishine T., Harada A.,

XA Konno H., Akiyama J., Nishi K., Kitsunai T., Tashiro H., Itoh M.,

XA Konno H., Akiyama J., Nishi K., Kitsunai T., Tashine T., Harada A.,

XA Konno H., Akiyama J., Nishi K., Kitsunai T., Tashine T., Kashiwagi K.,

XA Yamamoto R., Matsunoto H., Sakaguchi S., Ikagami T., Kashiwagi K.,

RA Fujiwake S., Inoue K., Toawa M., Ohara E., Watahiki M.,

XA Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,

RA Okazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y.;

"RIKEN integrated sequence analysis (RISA) system-384-format

RT sequencing pipeline with 384 multicapillary sequencer.";

RT Genome Res. 10:1757-1771(2000).
                                                                                                                                                                                                                                                                                                                                                                                Adachi J., Aizawa K., Akimura T., Arakawa T., Bono H., Carninci P., Fukuda S., Furuno M., Hanagaki T., Hara A., Hashizume W., Hayashida K., Hayatsu N., Hiramoto K., Hiraoka T., Hirozane T., A. Hayashida K., Hayatsu N., Hiramoto K., Hiraoka T., Hirozane T., Hara T., Kojima Y., Kondo S., Konno H., Kasukawa T., Katoh H., Kawai J., Kojima Y., Kondo S., Konno H., Kouda M., Koya S., A. Kurihara C., Matsuyama T., Miyazaki A., Murata M., Nakamura M., Nakamura K., Numazaki R., Ohno M., Ohsato N., Okazaki Y., A. Nishi K., Nomura K., Numazaki R., Ohno M., Ohsato N., Okazaki Y., A. Saitoh H., Sakai C., Sakai K., Sakazume N., Sano H., Sasaki D., Shibata K., Shinagawa A., Shiraki T., Sogabe Y., Tagami M. Tagawa A., Takahashi F., Takaku-Akahira S., Takeda Y., Tanaka T., Togami M., Togaru A., Toya T., Yasunishi A., Muramatsu M., Hayashizaki Y., Ramatu A., Toya T., Yasunishi A., Muramatsu M., Hayashizaki Y., Ramatu A., Kosoaja J., BaC34196.1; -
                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
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                                     Q6MI24;
Q6MI24;
05-JUL-2004
05-JUL-2004
05-JUL-2004
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STRAIN=C57BL/6J; TISSUE=Liver;
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STRAIN=C57BL/6J; TISSUE=Liver;
MEDLINE=20499374; PubMed=11042159;
05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein precursor (EC 3.6.3.14).
OrderedLocusNames=Bd3355;
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STRAIN=C57BL/6J; TISSUE=Liver;
MEDLINE=99279253; PubMed=10349636;
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"High-efficiency full-length cDNA cloning.";
Meth. Enzymol. 303:19-44(1999).
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STRAIN=C57BL/6J; TISSUE=Liver;
MEDLINE=21085660; PubMed=11217851;
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SEQUENCE FROM
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75.0%;
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Best Local :
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Q8MY38;
01-OCT-2002
01-OCT-2002
01-MAR-2004
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Sockett R.E., Schuster S.C.;
"A predator unmasked: life cy
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Science 303:689-692(2004).
EMBL; BX842655; CAE78158.1; -...
GO; GO:0016787; F:hydrolase activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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Profesia: Proteobacteria; Deltaproteobacteria; Bdellovibrionales;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SIGNAL
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SIGNAL 1 9 Potential.
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Science 303:689-692(2004).
EMBL; BX842655; CAE78158.1;
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Keller H., Lambert C., Evans K.J., (
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PubMed=14752164;
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Keller H., Lambert C.,
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Bdellovibrionaceae; Bdellovibrio.
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Pred. No. 2.8e
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Pred. No. 2.8e+02;
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A Lafontaine I., de Montigny J., Marck C., Neuveglise C., Talla B.,
A Goffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
A Boffard N., Frangeul L., Aigle M., Anthouard V., Babour A., Barbe V.,
A Boisrame A., Boyer J., Cattolico L., Confanioleri F., de Daruvar A.,
A Despons L., Fabre E., Fairhead C., Ferry-Dumazet H., Groppi A.,
A Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
A Hantraye F., Hennequin C., Jauniaux N., Joyet P., Kachouri R.,
A Nicaud J.M., Nikolski M., Oztas S., Ozier-Kalogeropulos O.,
A Nicaud J.M., Nikolski M., Oztas S., Ozier-Kalogeropulos O.,
A Pellenz S., Potter S., Richard G.F., Straub M.L., Suleau A.,
A Pellenz S., Potter S., Richard G.F., Straub M.L., Suleau A.,
A Swennene D., Tekaia F., Wesolowski-Louvel M., Westhof E., Wirth B.,
A Seniou-Meyer M., Zivanovic I., Bolotin-Fukuhara M., Thierry A.,
Bouchher C., Caudron B., Scarpelli C., Gaillardin C., Weissenbach J.,
Wincker P., Souciet J.L.,
"Genome arolution in vestra".
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Best Local
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Pfam; PP00098; zf-CCHC; 2.

PRINTS; PR00939; C2HCZNFINGER.

SMART; SM00343; ZnF C2HC; 2.

PROSITE; PS50158; ZF CCHC; 2.

PROSITE; PS50158; ZF CCHC; 2.
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Q6CJY8;
01-OCT-2004
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mol. Biol. Evol. 20:351-361(2003).
Mol. Biol. Evol. 20:351-361(2003).
Mol. Biol. Evol. 20:351-361(2003).
Mol. Biol. Evol. 20:351-361(2003).
GO, GO:0003676; F:nucleic acid bin
InterPro; IPR001878; Znf CCHC.
TEARNORS: Zf-CCHC; 2...
Genoscope;
Submitted (JUL-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; CR382126; CAG98459.1; -.
SEQUENCE 1036 AA; 118499 MW; 22DF98E5DBF3705C CRC64;
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Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Papilionoidea; Papilionidae; Papilioninae; Papilio.
NCBI_TaxID=66420;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kluyveromyces lactis (Yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Kluyveromyces.
                                                                               SEQUENCE FROM N.A.
STRAIN=NRRL Y-1140;
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STRAIN=NRRL Y-1140;
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Name=ORF1;
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(TrEMBLrel. 28, Last sequence update)
(TrEMBLrel. 28, Last sequence update)
Y-1140 chromosome F of strain NRRL Y-1140 of
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                                                                                                                                                   in yeasts.";
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Pred. No. 2.9e+02;
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Best Local S
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Best Local S
Matches 5
                                                                     Q91RA1
Q91RA1;
Q91RA1;
C91RA1;
C1-DEC-2001 (TrEMBLrel. 19, Cr
C1-DEC-2001 (TrEMBLrel. 26, La
C1-MAR-2004 (TrEMBLrel. 26, La
Core-spike protein lambda-2.
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01-JUL-1989
05-JUL-2004
                                                                                                                                                                                                                                                                                                           PIR; A28471; RMXRR3.

PIR; A28471; RMXRR3.

InterPro; IPR010311; Reovirus L2.

Pfam; PF06016; Reovirus L2; 1.

Pfam; PF06016; Reovirus L2; 1.

MRNA capping; mRNA processing; Nucleotidyltransferase; Transferase.

MRNA capping; mRNA processing; Nucleotidyltransferase; Transferase.

MRNA capping; MRNA processing; Nucleotidyltransferase; Transferase.

DOMAIN 941 968 Hydrophilic.

126 A297F0B8937018B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for comentities requires a license agreement (See http://www.isb-sib.ch/an or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=88058999; PubMed=2824487; Seliger L.S., Zheng K., Shatkin A.J.; Seliger Lucker and the sequence of recovirus L2 gene acid sequence of viral mRNA guanylyltransferase."; J. Biol. Chem. 262:16289-16293(1987).
                                                   Reovirus type 3 (strain Dearing) (T3D)
Viruses; dsRNA viruses; Reoviridae; On
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              guanylyltransferase).
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                                         Mammalian orthoreoviruses.
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  SEQUENCE
                       NCBI_TaxID=10886;
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Viruses; dsRNA viruses;
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CATALYTIC ACTIVITY: GTP + (5')PP-pur-mRNA = diphosphate
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(Rel. 11, Last sequence update)
(Rel. 44, Last annotation update)
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Reoviridae; Orthoreovirus
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Pred. No.
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Virology 287:33-348(2001).
EMBL; AP378010; AAK57514.1; -.
Interpro; IPR0010311; Recvirus L2.
Pfam; PF06016; Recvirus L2.
SEQUENCE 1289 AA; 143843 MW; E4
[1]
SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
STRAIN-B6.Sle1; TISSUE-Spleen;
STRAIN-B6.Sle1; TISSUE-Spleen;
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Z
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Z
Liang Z., Axie C., Chen C., Kreska D., Hsu K., Z
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Z
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Z
Liang Z., Xie C., Chen C., Kreska D., Hsu K., Z

"Antinuclear autoantibodies from B6.Sle1 mice."
Submitted (SEP-2003) to the EMBL/GenBank/DDBJ

"Av436911; AAR11071.1; -.
                                                                                                                                                                                                                            AAR11071 PRELIMINARY; PRT; 99 AA.
AAR11071;
02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
ANA immunoglobulin kappa light chain (Fragment).
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AAR11062;
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Luongo C.L., Nibert M.L.;
"Mammalian reovirus L2 gene and
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02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
ANA immunoglobulin kappa light chain (Fragment).
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Liang Z., Xie C., Chen C., Kre
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MEDLINE=21425393; PubMed=11531411;
T.A. Broering T.J., McCutcheon
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Mammalia; Eutheria;
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Rodentia;
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21-JUL-1986 (Rel. 01, Last seque
01-OCT-2004 (Rel. 45, Last annot
Ig kappa chain V-I region AU.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Ci
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                                                                                                                                                                              PROSITE; PS50835; IG_LIKE
                                                                                                                                                                                                   Pfam; PF00047; ig; 1.
SMART; SM00406; IGv;
                                                                                                                                                                                                                      PDB; 1JV5; X-ray; A=1-107.
GO; GO:0005576; C:extracellular; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                              Biophys.
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                                                                                                                                                                                                                                                                                                                                                                       Jones protein Au."
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MISCELLANEOUS: The Structure of the
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                                                                                        STRAIN=C57BL/6J; TISSUE-Head;
MEDLINE=20499374; PubMed=11042159;
Carninci P., Shibata Y., Hayateu N., Sugahara Y., Shiba
Konno H., Okazaki Y., Muramateu M., Hayashizaki Y.;
"Normalization and subtraction of cap-trapper-selected
prepare full-length cDNA libraries for rapid discovery
Genome Res. 10:1617-1630(2000).
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                                                                                                                                                                                                                                                                                                                                                                          the RIKEN Genome Exploration Research Group Phase I & II "Analysis of the mouse transcriptome based on functional 60,770 full-length cDNAs."; Nature 420:563-573(2002).
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STRAIN=C57BL/6J; TISSUE=Head;
MEDLINE=21085660; PubMed=11217851;
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STRALN=C57BL/6J; TISSUE=Head;
MEDLINE=99279253; PubMed=10349636;
Carninci P., Hayashizaki Y.;
"High-efficiency full-length cDNA cloning.";
meth. Enzymol. 303:19-44(1999).
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01-UN-2001 (TrEMBLrel. 17, Last annotation update)
Mus musculus 6 days neonate head cDNA, RIKEN full-length enriched
library, clone:5430437C04 product:hypothetical protein, full insert
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STRAIN=C57BL/6J; TISSUE=Head;
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Mammalia; Eutheria;
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RA Arakawa T., Bono H., Carninci P., Fukuda S., Fukunishi Y., Furuno M.,
RA Hanagaki T., Hara A., Hayatsu N., Hiramoto K., Hiraoka T., Hori F.,
RA Hanagaki T., Hara A., Hayatsu N., Hiramoto K., Hiraoka T., Hori F.,
RA Hanagaki T., Hara A., Hayatsu N., Izawa M., Kasukawa T., Kato H.,
Rawai J., Kojima Y., Konno H., Kouda M., Koya S., Kurihara C.,
RA Matsuyama T., Miyazaki A., Nishi K., Nomura K., Numazaki R., Ohno M.,
RA Matsuyama T., Okido T., Owa C., Saito H., Saito R., Sakai C., Sakai K.,
RA Okazaki Y., Okido T., Owa C., Saito H., Saito R., Sakai C., Sakai K.,
RA Sano H., Sasaki D., Shibata K., Shibata Y., Shinagawa A., Shiraki T.,
RA Sogabe Y., Suzuki H., Tagami M., Tagawa A., Takahashi F., Tanaka T.,
RA Sogabe Y., Toya T., Yamamura T., Yasunishi A., Yoshida K., Yoshino M.,
Muramatsu M., Hayashizaki Y.,
RA Shimited (AUG-2000) to the EMBL/GenBank/DDBJ databases.
REMBL; AKO19949; BAB31930.1; -.
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P., Konno H., Akiyama J., Nishi K., Kitsunai T., Tashiro H., Itoh M., Suni N., Tshii Y., Nakamura S., Hazama M., Nishine T., Harada A., Yamamoto R., Matsumoto H., Sakaguchi S., Ikegami T., Kashiwagi K., Fujiwake S., Inoue K., Togawa Y., Izawa M., Ohara E., Watahiki M., Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J., Okazaki Y., Muzamatsu M., Inoue Y., Kira A., Hayashizaki Y., "RIKEN integrated sequence analysis (RISA) system-384-format sequencing pipeline with 384 multicapillary sequencer.";
                                                                                                                                                                                                                                                                                                                                                                                                                          01-JAN-1988 (Rel. 06, Created)
01-APR-1988 (Rel. 07, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Ig kappa chain V-IV region B17 precursor.
Homo sapiens (Human)
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                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P06314;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hypothetical protein.
NON_TER 1
NON_TER 108 10
                                                                                                                                                      Submitted (OCT-1986) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                            Marsh
                                                                                                                                                                                                REVISION TO
                                                                                                                                                                                                                                                     "Detection of a unique human V kappa IV germline probe.";
                                                                                                                                                                                                                                                                                         MEDLINE=86041854; PubMed=2997713; Marsh P., Mills F., Gould H.;
                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KV4C HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
STRAIN=C57BL/6J; TISSUE=Head;
                                                                                                                                                                                                                                      Nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MGD; MGI:1925346; 5430437C04Rik.
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                                                                                                                                                                                                                                      Acids Res.
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                                                                                                                                                                                                                                      13:6531-6544(1985)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11561 MW;
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75.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8B2DBF861623608F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               134
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Q8VDD0;
Q1-MAR-2002 (TrEMBLrel. 20, C:
01-MAR-2002 (TrEMBLrel. 20, L:
01-MAR-2004 (TrEMBLrel. 26, L:
Anti-MOG Z12 variable light ol
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GO; GO:0005576; C:extracellular; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
Thierpro; IPR007110; Ig-like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Name=anti-MOG kappa;
Mus musculus (Mouse).
Eukaryota; Metazoa; Cr
Mammalia; Eutheria; Ro
                                                                                                                                                            Chernajovsky Y:;
submitted (CCT-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ416331; CACC94866.1; -.
PIR; G27887; G27887.
HSSP; P01834; 1MIM.
InterPro; IPR0037110; Ig-1ike.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IG; 1.
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 1.
Immunoglobulin V region; Sign
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SIGNAL
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STRAIN=BALB/c;
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                                                                                                              PROSITE; PS50835; IG_LIKE; 1.
NON_TER 134 134
SEQUENCE 134 AA; 14525 MW;
                                                                                                                                                                                                                                                                                                                                                  STRAIN=BALB/c
                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (JAN-2002)
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                                  66.1%;
ilarity 66.7%;
Conservative
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75.0%;
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Rodentia;
                                                                                                                14525 MW;
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                                  Score 37; DB Pred. No. 85; O; Mismatches
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Last annotation update)
chain (Fragment).
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Pred. No. 1
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Complementarity-determining-2.
Framework-3.
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Framework-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                CFDF8E2236E2D0CF CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6413A22FD0738832 CRC64;
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                                                                        DB 2;
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                                                                        Length 134;
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P SEQUENCE FROM N.A.

P SEQUENCE FROM N.A.

MEDLINE=99330421; PubMed=10403627;

XMEDLINE=99330421; PubMed=10403627;

LA Tohyama Y., Kasama-Yoshida H., Sakuma M., Kobayashi Y., Cao Ala Hasegawa M., Kojima H., Tamai Y., Tanokura M., Kurihara T.;

RT "Gene structure and amino acid sequence of Latimeria chalum (coelacanth) myelin DM20: phylogenetic relation of the fist Neurochem. Res. 24:867-873(1999).

RE MEL; AB02593; BAA84207.1; JOINED.

DR EMBL; AB025934; BAA84207.1; JOINED.

EMBL; AB025934; BAA84207.1; JOINED.

DR EMBL; AB025935; BAA84207.1; JOINED.

DR EMBL; AB025936; BAA84207.1; JOINED.
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Best Local S
Matches 6
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J. Bacteriol. 181:1585-1602(1999).
EMBL; AF079317; AAD04015.1; -.
PIK; T31291; T31291.
InterPro; IPR007110; Ig-like.
InterPro; IPR006311; Tat.
RTIGRFAMs; TIGR01409; TAT signal seq; 1.
PROSITE; PS50835; IG_LKE; 1.
PROSITE; PS50835; IG_LKE; 1.
PROSITE; PS50835; IG_LKE; 1.
POCENTIAL SIGNAL SIGN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q9PU09 PRELIMINARY;
Q9PU09;
01-MAY-2000 (TrEMBLrel. 1
01-MAY-2000 (TrEMBLrel. 1
01-JUN-2001 (TrEMBLrel. 1
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01-NOV-1998
01-NOV-1998
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                        Latimeria chalumnae (Latimeria) (Coelacanth).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Coelacanthiformes; Coelacanthidae; Latimeria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacteria; Proteobacteria; Alphaproteobacteria; Sphingomonadales;
Sphingomonadaceae; Novosphingobium.
NCBI_TaxID=48935;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hypothetical protein orf1251 precursor. Name=orf1251;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DM20 (Fragment).
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STRAIN=F199;
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(TrEMBLrel. 08, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
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75.0%;
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Pred. No. 1
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T.;
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Q6MQC7;
05-JUL-2004
05-JUL-2004
05-JUL-2004
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Q8UUT6;
01-MAR-2002
01-MAR-2002
01-MAR-2004
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NON_TER 1
SEQUENCE 242 AA; 2
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Pfam; PF01275; Myelin_PLP; 1.
PRINTS; PR00214; MYELINPLP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Werner H., Schweitzer J., Schachner M., Nave i
Submitted (DEC-2001) to the EMBL/GenBank/DDBJ
EMBL; AY070259; AAL59885.1; -
ZFIN; ZDB-GENE-001202-1; plp1a.
InterPro; IPR001614; Myelin PLP.
Pfam; PF01275; Myelin PLP; I.
PRINTS; PR00214; MYELINPLP.
SMART; SM00002; PLP; I.
Rendulic S., Jagtap P., Rosinus A.,
Keller H., Lambert C., Evans K.J.,
Sockett R.E., Schuster S.C.,
"A predator unmasked: life cycle of
                                                                                STRAIN=HD100 / DSM 50701 / ATCC 15356 / NCIB PubMed=14752164;
                                                                                                                                                                                   Bdellovibrio bacteriovorus.
Bacteria; Proteobacteria; Deltap
Bdellovibrionaceae; Bdellovibrio
                                                                                                                                                                                                                                             Hypothetical protein.
OrderedLocusNames=Bd0547;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cyprinidae; Dani,
NCBI_TaxID=7955;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Brachydanio rerio (Zebrafish)
Eukaryota; Metazoa; Chordata;
Actinopterygii, Neopterygii; T
Cyprinidae; Danio.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=plpla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DMalpha1
                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                             NCBI_TaxID=959;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PS01004; MYELIN PLP_2; 1.
245 AA; 27510 MW; 764B19BF8B949C9A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QYGVLPWT 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OYGLLPWT 178
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      242 AA;
                                                                                                                                                                                                                                                                                   (TrEMBLrel. 27, (TrEMBLrel. 27, (TrEMBLrel. 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
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75.0%;
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75.0%;
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                                                          Rosinus A.,
                                                                                                                                                                                                     Deltaproteobacteria; Bdellovibrionales;
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Pred. No. 1.5e
0; Mismatches
                                                                                                                                                                                                                                                                                      Last
Last
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Pred. No. 1.6e
0; Mismatches
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Last annotation update)
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annotation
    얁
                                        Goesmann A.,
    Bdellovibrio
                                                          Eppinger
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                                        , Baar C.,
Meyer F.,
    bacteriovorus
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RESULT 39
CAB78520
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RESULT QUESTION OF THE PROPERTY OF THE PROPERT
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Best Local S
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Best Local S
Matches 5
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01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Similarity to heat shock transcription factor.
Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eueurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Science 303:689-659(2004).
SCHENGE 303:689-659(2004).
EMBL; BX842647; CAB578520.1; -.
Complete proteome; Hypothetical protein.
SEQUENCE 330 AA; 37401 MW; E67D64135E164929
                                                                                                                        Kaneko T., Tabata S.;
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02-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genomic perspective.";
Science 303:689-692(2004).
EMBL; BX842647; CAE78520.1; -.
"Structural analysis of Arabidopsis thallana chromosome 5. features of the regions of 1,381,565 bp covered by twenty cphysically assigned P1 and TAC clones.";
DNA Res. 5:131-145(1998).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rendulic S., Jagrap P., Rosinus A.,
Keller H., Lambert C., Evans K.J.,
Sockett R.E., Schuster S.C.;
"A predator unmasked: life cycle of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bdellovibrio bacteriovorus.
Bacteria; Proteobacteria; Deltap
Bdellovibrionaceae; Bdellovibrio
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02-MAR-2004
                                                                                                                                                         MEDLINE=98344145;
Kaneko T., Kotani
                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                              NCBI_TaxID=3702;
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STRAIN=HD100 / DSM
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nes 5; Conserv
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                                                                                                                                                     Η.,
                                                                                                                                                         PubMed=9679202; H., Nakamura Y.,
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62.5%;
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27,
27,
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                          1,381,565 bp covered by twenty one TAC clones.";
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Pred. No.
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Pred. No. 2.1e+02;
2; Mismatches 1
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2; Mismatches
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                                                                                                                                                            Sato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  , Eppinger M., Baar
Goesmann A., Meyer
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DR EMBL; AB010693; BAB10879.1; -.

DR GO; GO;0003773; F:heat shock protein activity; IEA.

DR InterPro; IPR001816; F-box.

DR InterPro; IPR008546; FBD.

DR InterPro; IPR008945; Skp1_Skp2.

DR Pfam; PF00646; F-box; 1.

DR SMART; SM00579; FBD; 1.

DR PROSITE; PS50181; FBOX; 1.

NW Heat shock.

SQ SEQUENCE 388 AA; 45144 MW; 3117852CD27D24F5 CRC64;

Query Match

Best Local Similarity 75.0%; Pred. No. 2.5e+02;

Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

DD 256 HSYSKLGW 263

Search completed: December 17, 2004, 19:14:44

Job time: 28.8989 secs
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Copyright

GenCore version (c) 1993 - 2004

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Maximum Match 100%
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Perfect score:
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 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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length: 2000000000
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2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
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geneseqp2003as:*
geneseqp2003bs:*
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118.029 Million cell updates/
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AAB228151

AAB60401

AAB60401

AAB61586

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Modified
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Ganglioside

GD3 specific antibody related protein SEQ ID NO:

9

03-JUL-2001

(first

entry)

AAB81985;

AAB81985 standard;

protein;

119

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## ALIGNMENTS

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RESULT 1
AABB1985
ID AABB
XX A
                                                                               The present invention describes a monoclonal antibody which can react specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                             New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent
        Sequence 119 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-266143/27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-SEP-1999; 99JP-00278291.
06-APR-2000; 2000JP-00105088.
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                                                                                                                                                                                                                                                                                                                               Claim 20;
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                                                                                                                                                                                                                                                                                                                                                                                                               activity in cancer.
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                                                                                                                                                                                                                                                                                                                               Page 142-143; 183pp; Japanese
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Query Match
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Matches 119; Conserv

100.0%; ilarity 100.0%; Conservative 0

<u>,</u>

Mismatches

Score 624; DB 4; Pred. No. 8.2e-50;

Length 119; Indels

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Gaps

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Best Local Similarity
Matches 119; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention describes drugs contain a genetically-modified antibody against ganglioside GD3 or its fragment and at least 1 of a substance capable of activating immunocompetent cells and a substance having an antitumor activity in combination. The drugs can be used to treat tumour like melanoma and can provide a treatment with enhanced therapeutic effect and little side-reactions, particularly to relieve problems of side-effects during the conventional single administration. This sequence represents a protein associated with the anti- ganglioside GD3 antibody
AAB81987;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Drugs containing genetically-modified antibody against ganglioside GD3, its fragment, immunocompetent cell activators or/and antitumor agents i combination, applicable in treating malignant tumor like melanoma.
                                                         AAB81987 standard;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a monoclonal antibody which can react specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
  29-SEP-2000; 2000WO-JP006774.
                                                    05-APR-2001
                                                                                                                                                        Mus musculus.
                                                                                                                                                                                                                                 Ganglioside;
                                                                                                                                                                                                                                                                                 Ganglioside GD3 specific antibody related protein SEQ ID NO: 55.
                                                                                                                                                                                                                                                                                                                                     03-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                     AAB81989;
                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB81989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 41;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shitara K,
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                                                                                                                                                                                                                                 GD3; complementarity determining region; CDR; antibody;
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                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                    protein;
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Pred. No. 4.3e-49;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     119
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Matches 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors with low antigenicity, little side effects but potent activity in cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ganglioside GD3; anti-ganglioside GD3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modified ganglioside GD3 antibody associated protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABU11012;
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06-APR-2000;
                                                    Claim
                                                                                                    combination,
                                                                                                                                                                                                         WPI; 2003-067410/06.
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                                                                                                                                                                                                                                                                                                          (KYOW ) KYOWA HAKKO
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                                                                                                                         fragment,
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                                                                                            containing genetically-modified antibody against ganglioside GD3 ragment, immunocompetent cell activators or/and antitumor agents nation, applicable in treating malignant tumor like melanoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            109;
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                                                                                                                                                                                                                                                     Niwa R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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2000JP-00105088.
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                                          112-113; 121pp; Japanese.
                                                                                                                                                                                                                                                                                                          KOGYO KK
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Pred. No. 1.1e-45;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antibody; tumour; melanoma
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The invention describes drugs contain a

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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               against ganglioside GD3 or its fragment and at least 1 of a substance capable of activating immunocompetent cells and a substance having an antitumor activity in combination. The drugs can be used to treat tumo like melanoma and can provide a treatment with enhanced therapeutic effect and little side-reactions, particularly to relieve problems of side-effects during the conventional single administration. This seque represents a protein associated with the anti- ganglioside GD3 antibod
                                   The sequences given in AAR33256-57 represent rat heavy and light chain variable regions respectively. The DNA sequences encoding these proteins were used in the construction of humanised chimeric antibody expression vectors. In these humanised antibodies none of the amino acids of the nor human animal Ab variable region have been changed. (Updated on 25-MAR-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 119 AA;
                                                                                                               Claim
                                                                                                                                                  Humanised chimeric antibody prodn. against ganglioside GD3
                                                                                                                                                                                                               Shitara K,
                                                                                                                                                                                                                                                                 18-SEP-1991;
                                                                                                                                                                                                                                                                                         18-SEP-1992;
                                                                                                                                                                                                                                                                                                                  24-MAR-1993
                                                                                                                                                                                                                                                                                                                                           EP533199-A2
                                                                                                                                                                                                                                                                                                                                                                               Protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  humanised;
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12-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR33256;
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                                                                                                                                                                           AAQ33257
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                                                                                                              Page 29-30; 63pp;
                                                                                                                                       such as melanoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               variable; region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antibody; expression
                                                                                                                                                                                                               Hanai N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first en
                                                                                                                                                                                                                                                                 91JP-00238375
                                                                                                                                                                                                                                                                                         92EP-00116026.
                                                                                                                                                                                                                                                                                                                                                                                         /note= "Signal peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein;
                                                                                                                                                                                                                                        KOGYO CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92.8%;
91.6%;
                                                                                                                                                                                                               Hasegawa M,
                                                                                                                                                                                                                                                                                                                                                                  "Mature protein"
                                                                                                              English.
                                                                                                                                       neuroblastoma, etc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rat; immunoglobulin;
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Pred.
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No. 1.1e-45;
                                                                                                                                                                                                               Miyaji
                                                                                                                                                                                                                Kuwana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               heavy;
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Length 130;

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Best Local Sim
Matches 109;
                                                                                                         Query Match
Best Local S
Matches 109
                                                                                                                                                                                                                                                           chimeric human antibody H chain expression. mRNA from mouse anti-GD3 monoclonal Ab KM641-producing cells was isolated and KM641 H and L chain cDNAs isolated. The base sequences of the Ig variable regions in KM641 H chain cDNA (pKM641HA3) and KM641 L chain cDNA (pKM641LA2) are given in AAQ45438-39. A KM641-derived chimeric human Ab H chain expression vector was constructed by joining the H chain variable region gene from pKM641HA3 to a vector for chimeric human Ab H chain expression using the synthetic DNAs given in AAQ63439 and AAQ63440
                                                                                                                                                                                                                  Sequence 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Humanised antibody specific for ganglioside GM2 - used for producing a cytocidal effect on cancers such as melanoma, neuroblastoma and glioma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2 describes the construction of the vector pChi641HA1 for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Page 116-117; 191pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nakamura K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-SEP-1992;
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                                                                                                           109;
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                                                                                                      h 92.8%;
Similarity 91.6%;
09; Conservative
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EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                   EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Koike M,
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1. .10
/label= sig_peptide
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••
                                                                                                      Score 579; DB 2;
Pred. No. 1.2e-45;
5; Mismatches 5
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Pred. No. 1.2e-45;
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                                                                                                                                                        Length 130;
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                                                                                                                                                                                                                  Query Match
Best Local S
Matches 109
                                                                                                                                                                                                                                                                                                  This immunoglobulin region was isolated from pKM641HA3. This sequence has no methionine initiation codon and the leader sequence was partly lacking. The chimeric human antibodies are useful in the treatment of cancer, especially that which is of neural ectodermal origin. In contrast to prior art constructs based on mouse monoclonal antibodies, the chimeric human antibodies do not cause anti-mouse immunoglobulin production. The chimeric human antibodies have a prolonged half-life and a reduced frequency of adverse effects when compared to mouse monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus
                                     AAB01627;
                                                           AAB01627
                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Col 99-101; 188pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Chimeric
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUN-1995;
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light chain; amino acid.
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                                                                                                                                                                               1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                                                     SDSVXGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS
                                                                                                                                                                    EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
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                                                           standard;
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                                                          protein; 130
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91.6%;
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Pred. No. 1.2e-45;
5; Mismatches 5;
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                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is a murine immunoglobulin heavy chain variable region from plasmid KM-641. The coding sequence was used in the creation of an expression vector, along with the sequence for a human antibody, to produce humanised chimaeric antibodies, which can be used to treat cancer. Humanised chimaeric antibodies are more effective than mouse antibodies as they do not provoke a reaction in the human and side effects, such as the formation of anti-mouse immunoglobulin antibody and the rapid half-life of the immunoglobulins, do not occur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New humanized chimera antibody KM-871 useful for treating cancer, comprises variable region of mouse monoclonal antibody, reactive with ganglioside and human antibody constant region.
                                                                                                                                                                                                                                                                                                                                                                                Sequence 130 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Protein
Mus musculus.
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18-SEP-1992;
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                                       Ganglioside;
                                                                  Ganglioside
                                                                                            03-JUL-2001
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                                                                                                                                               AAB81977 standard; protein; 138 AA
                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                            109;
                                                                                                                                                                                                                                           61
                                                                                                                                                                                                                    71
                                                                                                                                                                                                                                                                       H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunoglobulin; H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunoglobulin heavy chain variable region.
                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                      EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119
                                                                                                                                                                                                                                                                       EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                   G
B
3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hanai N,
                                                                                                                                                                                                                                                                                                                            Conservative
                                        GD3;
                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            91JP-00238375
92EP-00116026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99EP-00124345
                                                                specific antibody related protein SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /label= signal_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label= mature_immunoglobulin_heavy_chain_variable region
                                       complementarity determining
                                                                                                                                                                                                                                                                                                                                        92.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hasegawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chain; heavy chain; variable region; cancer;
                                                                                                                                                                                                                                                                                                                                         Score 579; DB 3; Pred. No. 1.2e-45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Σ,
                                                                                                                                                                                                                                                                                                                            Mismatches
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                                       region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kuwana
                                                                                                                                                                                                                                                                                                                                                    Length 130;
                                                                                                                                                                                                                                                                                                                            Indels
                                        CDR; antibody;
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RESULT 11
ABUIIO02
ID ABUII
XX ABUIII
XX ABUIII
XX ABUIII
XX GANGII
CX GANGII
XX GANGI
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Best Local Simi
Matches 109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 138 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a monoclonal antibody which can react specifically with ganglioside GD3. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent
                                                                                                         Shitara K,
                                                                                                                                                                                                               29-MAR-2001; 2001JP-00097483
                                                                                                                                                                                                                                                                    29-MAR-2002; 2002WO-JP003170
                                                                                                                                                                                                                                                                                                                        10-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                             WO200278739-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ganglioside GD3; anti-ganglioside GD3 antibody; tumour; melanoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Modified ganglioside GD3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABU11002 standard; protein; 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 138-139; 183pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                activity in cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (KYOW ) .KYOWA HAKKO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-SEP-1999; 99JP-00278291
06-APR-2000; 2000JP-00105088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-SEP-2000; 2000WO-JP006774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200123432-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the invention
                                                       2003-067410/06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                KYOWA HAKKO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS
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                                                                                                      Niwa R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                KOGYO KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
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                                                                                                            Kanazawa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antibody associated protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 579; DB 4; 1
Pred. No. 1.3e-45;
5; Mismatches 5;
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                                                                                                            Asada
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 138;
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Drugs containing genetically-modified antibody against ganglioside

GD3

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RESULT 12
AAB811991
ID AAB81
XX AAB811
XX AAB81
XX AAB81
XX AAB81
XX AAB81
XX AAB81
XX AAB81
XX Gangl

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention describes drugs contain a genetically-modified antibody against ganglioside GD3 or its fragment and at least 1 of a substance capable of activating immunocompetent cells and a substance having an antitumor activity in combination. The drugs can be used to treat tumour like melanoma and can provide a treatment with enhanced therapeutic effect and little side-reaction, particularly to relieve problems of side-effects during the conventional single administration. This sequence represents a protein associated with the anti- ganglioside GD3 antibody
                                                                                  The present invention describes a monoclonal antibody which can react specifically with ganglioside GOJ. The antibody and its derivatives are useful in the diagnosis and therapy of tumours, particularly cancer diagnosis. The present sequence is a protein used in the exemplification
                                                                                                                                                                                                                                                                                                                              New human type complementation-determining region-transplanted antibody and derivatives against ganglioside GD3, useful in diagnosis and therapy of e.g. tumors, with low antigenicity, little side effects but potent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       its fragment, immunocompetent cell activators or/and antitumor agents in combination, applicable in treating malignant tumor like melanoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-SEP-1999; 99JP-00278291
06-APR-2000; 2000JP-00105088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ganglioside
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 138
                                                                                                                                                                                                                                                  Claim
                                                                                                                                                                                                                                                                                                           activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hanai N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-SEP-2000; 2000WO-JP006774
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (KYOW ) KYOWA HAKKO KOGYO KK
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                                                                  invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                           in cancer.
                                                                                                                                                                                                                                            Page 175-179; 183pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shitara K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GD3 specific antibody related protein SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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ج
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91.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nakamura
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Pred. No. 1.3e-45;
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RESULT 13
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XX EP138
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                          CC that binds to melanoma cell adhesion molecules (MCAM), comprising a compression and comprising and compression and comprising the melanoma cell adhesion molecules (MCAM), comprising a comprise and a cids as given in the specification. The invention comprises: a bioconjugate comprising the melanoma cell adhesion comprises: a bioconjugate comprising the melanoma cell adhesion comprises: a bioconjugate comprising the protein and/or the bioconjugate and a container; a pharmaceutical composition comprising compressing the protein; a vector comprising the nucleic acid compression to prevention in the manufacture of a medicament for the concurring cancer cells, where invasion and/or metastasis of naturally comprising cancer cells depends on melanoma cell adhesion molecule (MCAM) compositions have cytostatic activity. The MCAM binding protein and compositions have cytostatic activity. The MCAM binding protein and compositions have cytostatic activity. The MCAM binding protein and bioconjugate are useful for the detection of MCAM, or for identifying a metastasis in a patient, involves administering the MCAM binding protein and bioconjugate are useful for the MCAM mediate invasiveness and/or metastatic control of the manufacture of the MCAM binding protein and bioconjugate are useful for the detection of MCAM, or for identifying a metastasis in a patient, involves administering the MCAM binding protein and bioconjugate are useful for treating or preventing the MCAM mediated invasiveness and/or metastatic control of the model of the metastatic control of the metastatic control of the MCAM binding protein and bioconjugate are useful for treating or preventing the MCAM binding protein and bioconjugate are useful for treating or preventing the MCAM binding protein and bioconjugate are useful for treating the MCAM binding protein and bioconjugate are useful for treating the MCAM binding protein and bioconjugate are useful for treating the MCAM binding protein and bioconjugate are useful for treating the MCAM binding protei
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Best Local Sim:
Matches 109;
   metastasis in a patient, involves administ and bioconjugate to inhibit MCAM mediated potential. The MCAM inhibiting molecule is
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel polypeptide useful for detecting melanoma cell adhesion molecules for treating or preventing metastasis in patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; ADO25162
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Melanoma cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; SEQ ID NO 5; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61
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Pred. No. 6e-45;
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s useful for inhibiting
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RESULT 14
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Matches 94
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; interferon alpha; IFNalpha; insulin-dependent diabetes mellitus; autoimmune disease; systemic lupus erythematosus; autoimmune thyroiditis; antibody therapy; immunosuppressive; antiinflammatory; dermatological;
                                                                                                                                                                                                                                                                                                                                                                                    22-FEB-2001; 2001US-0270775P 09-JAN-2002; 2002US-00044896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-JAN-2002; 2002WO-US002709.
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                                                                                         New anti-interferon alpha monoclonal antibody, 9F3, useful for treating autoimmune diseases e.g. insulin-dependent diabetes mellitus, systemic
                                                                                                                                                                                                                                                  Chuntarapai A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Region
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                                                                                                                                                                                                                                                         Kim JK,
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                                                           e.g. insulin-dependent and autoimmune thyroid:
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                                                                                                                                                                                                                                                         Presta LG,
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Pred. No. 3.4e-38;
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ErbB2 receptor; anti-ErbB2 antibody; tumour;

cancer; breast cancer;

leukemia;

cytostatic;

carcinoma.

Human Vh consensus framework humIII.

29-MAY-2001

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entry)

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                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to novel anti-interferon alpha (IFNalpha) monoclonal antibody, 973 which binds to and neutralises a biological activity of IFN-alpha subtypes. The anti-IFNalpha monoclonal antibodies of the invention are useful for treating autoimmune diseases such as insulin-dependent diabetes mellitus, systemic lupus erythematosus and autoimmune thyroiditis. They are useful as reagents in diagnostic assays for IFN-alpha expression, for the affinity purification of various IFN-alpha subtypes from recombinant cell culture or natural resources and for the detection of IFN-alpha in diagnostic assay methods. Sequences of the invention are also useful in antibody therapy. The present sequence is
AAB62088
                                     AAB62088 standard;
                                                                                                                                                                                                                                                                                                                                                                              Sequence 119 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                      human consensus heavy chain variable domain subgroup III (huIII)
                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                       61
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                                                                                                                                                                                                                                                                                                   96;
                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                            SDSVKGRFTISRDNSKNTLYLOMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119
                                                                                                                                                                                                                                             EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
                                                                                                                                     ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARGRVG--YYDYWGQGTLVTVSS 117
                                                                                                                                                                                                                    EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVAVISGDGGSTYY
                                                                                                                                                                                                                                                                                                   Conservative
                                   protein;
                                                                                                                                                                                                                                                                                                                    80.0%;
                                       119
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                                                                                                                                                                                                                                                                                               Score 499; DB 5;
Pred. No. 2.6e-38;
8; Mismatches 13
                                       $
                                                                                                                                                                                                                                                                                                   13;
                                                                                                                                                                                                                                                                                                                                         Length
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ARESULT 15
ARABEZ
ARA ARABEZ
ARA ARABEZ
XX XARABEZ
XX ARABEZ
XX ZARABEZ
XX ZARABEZ
XX ExbBZ
XX ExbBZ
XX ExbBZ
XX Human
X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Treating a human patient with a disorder characterized by overexpression of ErbB2 receptor such as a tumor or cancer (e.g. metastatic breast carcinoma) comprises administering several dosages of an anti-ErbB2
                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-235058/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Baughman SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-AUG-1999; 99US-0151018P.
23-JUN-2000; 2000US-0213822P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-MAR-2001.
                                                                                                                                                                                                                                                                                   The invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200115730-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shak S;
                                                                                                                                                                                                                                                                           relates to treating cancer
                                                                                                                                                                                                                                                                                                                                                                 5B;
                                                                                                                                                                                                                                                                                                                                                         71pp; English
                                                                                                                                                                                                                                                                               in a human susceptible
                                                                                                                                                                                                                                                                                   or
P
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Disclosure; Page 98; 100pp;

English

diagnosed with a disorder characterized by overexpression of ErbB2 receptor. The method comprises administering an initial dose of at least approx. S mg/kg of the anti-ErbB2 antibody, and administering subsequent doses of the antibody in approximately the same amount or less than the initial dose. The method is useful for treating a disorder characterized by overexpression of ErbB2 receptor, particularly a benign or malignant

The cancer

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RESULT 16
AAB60401
ID AAB60
XX AAB60
XX AAB60
XX AAC1
XW AAC1
KW AAC1
KW AAC1
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KW COLox
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Matches
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The invention relates to a method for treating cancer in a human patient, wherein the cancer expresses epidermal growth factor receptor (EGFR), comprising administering an antibody which binds ErbB2 (HER2; AAB60408). In particular, the anti-ErbB2 antibody is the murine monoclonal antibody 2C4 (AAB60396, AAB60397) or a humanised version of 2C4 (AAB60398, AAB60399). The invention also encompasses an isolated nucleic acid encoding a humanised ErbB2-binding antibody; vectors and host cells comprising such nucleic acids; the recombinant production of a humanised ErbB2-binding antibody; and an immunoconjugate comprising a humanised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer, endometrial carcinoma, salivary gland carcinoma, kidney cancer, liver cancer, prostate cancer, vulval cancer, thyroid cancer, hepatic carcinoma and various types of head and neck cancer. The present sequence represents the human variable heavy chain consensus framework humili
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer, small-cell lung cancer, non-small cell lung cancer, pancreatic cancer, gastrointestinal cancer, glioblastoma, cervical cancer, ovarian cancer, liver cancer, bladder cancer, hepatoma, colon cancer, colorectal
                                                                                                                                                                                                                                                                                                                                                                                                       Treating cancer factor receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    heavy chain variable region; cancer; cytostatic; EGFR-expressing epidermal growth factor receptor; colon cancer; rectal cancer; tu colorectal cancer; non-small cell lung cancer; metastatic breast affinity purification; subgroup III framework.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 119 AA;
                                                                                                                                                                                                                                                                                                                 Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-080862/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200100245-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-ErbB2 monoclonal antibody 2C4; HER2; humanised; VH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Consensus human heavy chain subgroup III framework,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB60401 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GETH ) GENENTECH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (heavy subgroup III)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       particularly metastatic breast carcinoma), leukemia, squamous cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARGRVGYSLYDYWGQGTLVTVSS
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                                                                                                                                                                                                                                                                                                                 Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Presta LG,
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                                                                                                                                                                                                                                                                                                              7B; 89pp;
                                                                                                                                                                                                                                                                                                                                                                                                 in a human, where the cancer expresses epidermal growth (EGFR), comprises administering an antibody which binds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein;
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79.8%;
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                                                                                                                                                                                                                                                                                                              English.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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Best Local S
Matches 95
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ErbB2-binding antibody and a cytotoxic drug. The ErbB2-binding antibodic act by antagonising ErbB receptors, and as inhibitors of transforming growth factor alpha (TGF-alpha)-activated mitogen activated protein kinase (MAPK). The method of the invention is used for treating cancer, especially colon cancer, rectal cancer, colorectal cancer, lung cancer (especially non-small cell lung cancer), or breast cancer (especially morematic cancer). The antibodies may also have non-therapeutic
                                                                                                                                                                                                                                                               Sequence 119 AA;
61
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                                                                                                                                                                        95;
                                                                                                                                                                                               Similarity
                                                                                                                           EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGRVGYSLYDYWGQGTLVTVSS
                                                                                    EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVAVISGDGGSTYY
                                                                                                                                                                        Conservative
                                                                                                                                                                                             79.8%;
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                                                                                                                                                                                             Score 498; DB 4;
Pred. No. 3.3e-38;
                                                                                                                                                                             Mismatches
                                                                                                                                                                             16;
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RESULT 17
AAB61586
ID AAB61586
AC AAB61
XX AAB61
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XX AAB61
XX Human
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XX Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human variable heavy chain consensus sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-APR-2001
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SLOAN KETTERING INST CANCER RES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sliwkowski
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The BrbB family of receptor tyrosine kinases are important mediators of cell growth, differentiation and survival. The receptor family includes four distinct members including Epidermal Growth Factor Receptor (EGFR or ErbB1), HER2 (ErbB2 or p185^neu), HER3 (ErbB3) and Her4 (ErbB4 or tyro2). The present invention relates to a method for treating prostate cancer. The method comprises administering an antibody which binds ErbB2 and blocks ligand activation of an ErbB receptor. Preferably, the antibody

Example 3;

Fig 7; 93pp;

English

Treating prostate cancer in a human comprises administering an antibody which binds ErbB2 and blocks ligand activation of an ErbB receptor.

TGF-

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RESULT 18
AAU74541
ID AAU74
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Best Local
The invention relates to treating a tumour in a mammal, where the tumour is characterised by the overexpression of an epidermal growth factor receptor (ErbB) and does not respond or respond poorly, to treatment with an anti-ErbB antibody, comprising administering to the mammal an anti-ErbB antibody-maytansinoid conjugate. The method is useful for treating cancer or tumours of the breast, ovary, stomach, endometrium, salivary gland, lung, kidney, colon, colorectum, thyroid, pancreas, prostate and bladder, preferably breast cancer. The breast cancer is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  anti-ErbB antibody-maytansinoid conjugate; cancer; tumour; breast; ovastomach; endometrium; salivary gland; lung; kidney; colon; colorectum; thyroid; pancreas; prostate; bladder; ErbB2; neuronal disorder; antibody; glial disorder; astrocytal disorder; hypothalamic disorder; antibody; glandular disorder; macrophagal disorder; pythelial disorder; stromal disorder; blastocoelic disorder; inflammatory disorder; angiogenic disorder; immunological disorder; heavy chain variable reginancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     blocks binding of monoclonal antibody 2C4 to ErbB2 and/or blocks alpha activation of mitogen-activated protein kinase (MAPK). The sequence is the consensus framework for the human variable heavy This sequence was used in a sequence alignment homology with the heavy chain of antibody 2C4 (see AAB61582)
                                                                                                                                                                                                                                                                                                 (ERIC/)
(SCHW/)
(SLIW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; subgroup V\_H III; ErbB; epidermal growth factor receptor; anti-ErbB antibody-maytanainoid conjugate; cancer; tumour; breas
                                                                                                                                            Claim
                                                                                                                                                                                                                                       WPI; 2002-163686/21.
                                                                                                                                                                                                                                                                    Erickson
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                                                                                                                                                                        antibody-maytansinoid
                                                                                                                                                                                                                                                                                                                                                                             16-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                            16-MAR-2001; 2001US-00811123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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                                                                                                                                                                                                                                                                                                 SCHWALL R.
SLIWKOWSKI M.
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                                                                                                                                                                                     tumor characterized by ove, ErbB or cancer in mammal,
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2000US-0238327P
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                                                                                                                                                                                                                                                                  Schwall
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                                                                                                                                         1; 93pp;
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                                                                                                                                                                      conjugate to
                                                                                                                                         English.
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Pred. No. 3.3e-38;
8; Mismatches 16
                                                                                                                                                                                      overexpression of epidermal mal, comprises administering
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                                                                                                                                                                      the mammal.
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                                                                                                                                                                                      growth factor
anti-ErbB
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chain.
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RESULT 19
ADE71455
ID ADE71
XX Cytos
KW Crbs
KW ErbB2
KW Adeno
KW Adeno
KW Adeno
KW Erbma
KW Cervi
KW Adeno
KW Cervi
KW Adeno
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KW Adeno
KW Adeno
KW ADE71
KW ADE71
XX US200
XX 10-OC
XX 25-JU
PR 23-JU
XX ITEST
PT Treat
PT Sliwk
XX Treat
PT Treat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KW cytostatic; ErbB2 antagonist; tyrosine kinase inhibitor; cancer; ErbB2;
KW ErbB2 antibody; carcinoma; lymphoma; blastoma; sarcoma; liposarcoma;
KW neuroendocrine tumour; mesothelioma; schwanoma; meningioma;
KW adenocarcinoma; melanoma; leukaemia; lymphoid malignancy;
KW squamous cell cancer; epithelial squamous cell cancer; lung
kW small-cell lung cancer; non-small cell lung cancer;
KW small-cell lung cancer; non-small cell lung cancer;
KW cancer of the peritoneum; hepatocellular cancer; gastric cancer;
KW cancer of the peritoneum; hepatocellular cancer; gastric cancer; glioblastoma;
KW cervical cancer; ovarian cancer; liver cancer; bladder cancer; hepatoma;
KW cervical cancer; rectal cancer; colorectal cancer;
KW endometrial carcinoma; uterine carcinoma; salivary gland carcinoma;
KW kidney cancer; renal cancer; prostate cancer; vulval cancer;
KW kidney cancer; hepatic carcinoma; anal carcinoma; penile carcinoma;
KW testicular cancer; oesophageat cancer; tumour of the biliary tract;
KW head and neck cancer, human antibada; variatical cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ঠ
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Best Local (
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                                                                                                                                                                                                 WPI; 2004-020226/02.
                                                                                                                                                                                                                                                  Sliwkowski MX
                                                                                                                                                                                                                                                                                                                                              25-JUN-1999; 99US-0141316P.
23-JUN-2000; 2000US-00602812.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human antibody variable heavy chain framework.
                                                                                                                                                                                                                                                                                                (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and neck cancer; human antibody; variable heavy chain
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79.8%;
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Pred.
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No. 3.3e-38;
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Treating cancer (e.g. carcinoma, lymphoma or sarcoma) that exerbs by administering to a patient an anti-Erbs2 antibody, experionally an epidermal growth factor receptor-targeted drug

lymphoma or sarcoma) that expresses

or

Example 3; SEQ ID NO tyrosine kinase

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56pp; English

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RESULT 20
ADJ889
ID ADJ888
XX ADJ88
XX ADJ88
XX ADJ88
XX ADJ88
XX ADJ88
CT 06-Ma
XX Cance
XX Cance
XX Unide
XX Unide
XX Unide
XX Unide
XX 22-Ja
PP 22-Ju
PR 23-Ju
PR 23-Ju
PR 23-Ju
PR 23-Ju
PR 23-Ju
PR 10-OC
XX WPI;
XX WPI;
XX WPI;
XX WPI;
XX WPI;
YX Treat
PT Colox
PS Examp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             particularly a human. The cancer includes carcinoma, lymphoma, blastoma, carcoma, liposarcoma, neuroendocrine tumour, mesothelioma, schwanoma, meningioma, adenocarcinoma, melanoma, leukaemia, lymphoid malignancy, captuamous cell cancer, epithelial squamous cell cancer, lung cancer, small cell lung cancer, non-small cell lung cancer adenocarcinoma of the lung, cancer of the peritoneum, hepatocellular cancer, gastric or stomach cancer, gastrointestinal cancer, panoreatic cancer, glioblastoma, cervical cancer, ovarian cancer, liver cancer, bladder cancer, phopatoma, breast cancer, colon cancer, cretal cancer, colorectal cancer, endometrial or uterine carcinoma, salivary gland carcinoma, kidney or renal cancer, prostate cancer, vulval cancer, thyroid cancer, hepatic carcinoma, anal carcinoma, penile cancinoma, tract, or head and neck cancer. This is the amino acid sequence of human antibody variable heavy chain framework that was used to produce a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 95
                                                                                                                                                                                                                                                        25-JUN-1999; 99US-0141316P.
23-JUN-2000; 2000US-00602812.
10-OCT-2002; 2002US-00268501.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention describes a method of treating cancer that expresses comprising administering to a patient an antibody that binds ErbB2. Specifically claimed are antibodies that bind ErbB2, particularly monoclonal antibody 2C4 or humanised 2C4, or monoclonal antibody 4D
                                                           Treating cancer that expresses ErbB2 e.g., colorectal cancer comprises administering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human variable heavy chain consensus peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 119
                                                                                                                                WPI; 2004-121529/12
                                                                                                                                                                                                                                                                                                                                            27-JUN-2003; 2003US-00608626
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cancer; ErbB2; gene therapy; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-MAY-2004
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                                                                                                                                                                                                                  (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    humanised antibody.
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                                                                                                                                                                      SM,
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95; Conservative
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                                                                                                                                                                        Sliwkowski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The methods are useful for treating cancer in a patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          79.8%;
79.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 498; DB 8; Pred. No. 3.3e-38; 8; Mismatches 16;
                                                              an antibody that binds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 119;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                ErbB2
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Example 3;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          carcinoma, lymphoma, blastoma, medulloblastoma, retinoblastoma, sarcoma, carcinoma, synovial cell sarcoma, neuroendocrine tumour, carcinoid tumour, gastrinoma, islet cell cancer, mesothelioma, schwannoma, acoustic neuroma, meningioma, adenocarcinoma, melanoma, leukaemia, lymphoid malignancy, squamous cell cancer, epithelial squamous cell cancer, lung cancer, small-cell lung cancer, non-small cell lung cancer, lung cancer of the lung, squamous carcinoma of the lung, squamous carcinoma of the lung, squamous carcinoma of the lung, cancer of the peritoneum, hepatocellular cancer, gastric or stomach cancer, cancer of the cancer, ovarian cancer, liver cancer, bladder cancer, hepatoma, breast cancer, colon cancer, rectal cancer, colorectal cancer, endometrial or uterine carcinoma, salivary gland carcinoma, kidney or renal cancer, prostate cancer, vulvar cancer, thyroid cancer, hepatic carcinoma, anal carcinoma, penile carcinoma, testicular cancer, oesophageal cancer, a tumour of the biliary tract or head and neck cancer with anti-ErbB2 antibodies. The invention is also useful in gene therapy. The present sequence is human variable heavy chain consensus peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local (
The present invention relates to identifying a tumor responsive to treatment with an anti-HER2 antibody involves detecting the present an HER2/HER3 and/or HER2/HER1 protein complex in a sample of the t
                                                                                                          Identifying a tumor responsive to treatment wit useful in treating cancer, comprises detecting HER2/HER3 and/or HER2/HER1 protein complex.
                                                                                                                                                                                       WPI; 2004-156546/15.
                                                                                                                                                                                                                                                                                                    15-JUL-2002; 2002US-0396290P
20-JUN-2003; 2003US-0480043P
                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2004008099-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tumor; anti-HER2 antibody; HER2/HER3; HER2/HER1; Cytostatic; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Variable heavy chain consensus sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADN12055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADN12055 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present
                                                                         Disclosure; SEQ ID NO 6; 105pp; English.
                                                                                                                                                                                                                          Koll H,
                                                                                                                                                                                                                                                                                                                                                          11-JUL-2003; 2003WO-US021590
                                                                                                                                                                                                                                                                 (GETH )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Н
                                                                                                                                                                                                                                                                 GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                            Bossenmaier B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARGRVGYSLYDYWGQGTLVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EVOLVESGGGLVOPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVAVISGDGGSTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention relates to methods for treating cancer such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein; 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      79.8%;
79.8%;
                                                                                                                                                                                                                            Mueller H,
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Pred. No. 3.3e-38;
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                                                                                                                                                                                                                            Sliwkowski MX,
                                                                                                                                                     with
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                                                                                                                                 presence
                                                                                                                                                     anti-HER2 antibody,
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                                                                                                                                     of a
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RESULT 22
ADP43329
ID ADP43
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Best Local
                         This invention refers to a novel method for treating a non-malignant disease that involves abnormal activation or production of an epidermal growth factor receptor (BrbB) receptor or an ErbB ligand. Specifically, it refers to the use of humanised murine anti-ErbB2 antibodies to block ligand activation of the ErbB receptor, where the ErbB ligand is a transforming growth factor alpha that promotes mitogen-activated proteir kinase (MAPK). The present invention describes conjugating the monoclona murine antibody 2C4, or more particularly the humanised antibody 574, to
                                                                                                                                                                                                                                                                                                                                                                          Treating non-malignant disease or disorder such as psoriasis, endometriosis, involving abnormal activation or production of epidern growth factor receptor or ErbB ligand by administering antibody that binds ErbB2 to mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gynaecological; dermatological; vasotropic; antiarteriosclerotic; cardiant; antianginal; antilipaemic; hypotensive; antiasthmatic; antiallergic; antiinflammatory; antitussive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            epidermal growth factor receptor; ErbB; ErbB2;
transforming growth factor alpha; benign hyperproliferative disorder;
psoriasis; endometriosis; scleroderma; vascular disease; atherosclero;
restenosis; colon polyps; fibroadenoma; respiratory disease;
chronic bronchitis; cystic fibrosis; cytostatic; antipsoriatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sliwkowski MX,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADP43329 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The method is useful in identifying a tumor as responsive to treatment with an anti-HER2 antibody. The methods and antibodies are useful in treating a condition or disorder including tumor or cancer, e.g. breast, prostate, lung, colorectal or ovarian cancer. The present sequence
                                                                                                                                                                                                                                                                                                                Example 3; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-NOV-2002; 2002US-0428027P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-NOV-2003; 2003WO-US037367
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human; variable heavy kappa chain; antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human monoclonal variable heavy kappa chain antibody SeqID 6.
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   cytotoxic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2004-450361/42.
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   therapeutic agent such that
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79.8%;
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Pred. No. 3.3e-38;
8; Mismatches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disease; atherosclerosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    epidermal
                                                             monoclonal
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Best Local (
                                                                                                                                            Matches
                                                                                                                                                                                                                                              benign hyperproliferative disorder, psoriasis, endometriosis, scleroderma, vascular disease (such as atherosclerosis or restenosis) colon polyps, fibroadenoma or respiratory disease (such as chronic bronchitis or cystic fibrosis). Accordingly, they exhibit cytostatic, antipsoriatic, gynaecological, dermatological, vasotropic, antiarteriosclerotic, cardiant, antianginal, antilipaemic, hypotensiv antiasthmatic, antiallergic, antianflamatory and antitussive activit This polypeptide sequence is the human variable heavy kappa chain pro used to humanise the murine anti-ErbB2 proteins of the invention.
                                                                                                                                                                                                                Sequence 119
61
                                 61
                                                                     -
                                                                                             1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
                                                                                                                                          95;
                                                                                                                                                           Similarity
               SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS
                                                                   EVOLVESGGGLVOPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVAVISGDGGSTYY
ADSVKGRFT1SRDNSKNTLYLQMNSLRAEDTAVYYCARGRVGYSLYDYWGQGTLVTVSS
                                                                                                                                         Conservative
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                                                                                                                                                           79.8%;
                                                                                                                                         Score 498; DB Pred. No. 3.3e 8; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                              as atherosclerosis or restenosis),
                                                                                                                                                        DB 8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         idiopathic pulmonary fibrosis; benign prostate hypertrophy; colon cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nephritis; glomerulosclerotic renal disease; lung cancer
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24-APR-2002; 2002WO-US012801

24-APR-2001; 2001US-0285683P

(FARB ) (MORP-) BAYER CORP ద్ద

N-PSDB; 2003-129114/12. ABZ74782

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Knorr AM, Schauer

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Hirth-Dietrich

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New human anti-TIMP-1 (tissue inhibitor of metalloprotease-1) antibodies, for diagnosing or ameliorating the symptoms of a disorder in which TIMP-1 is elevated, e.g. liver fibrosis, benign prostate hypertrophy or lung TIMP-1

Claim 20; Page 153-154; 228pp; English.

The invention relates to a novel purified preparation of a human antibody, which binds to a tissue inhibitor of metalloprotease-1 (TIME and neutralises a matrix metalloprotease (MMP)-inhibiting activity of -1. The antibody comprises a variable heavy chain (VHC) DR3 region and variable light chain (VLC) DR3 region. An antibody preparation of the invention has hepatotropic, cytostatic, nephrotropic (TIMP-1)

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Loehning C,
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                                                                                      New antibody molecule capable of specifically recognizing two regions the beta-A4 peptide, useful for diagnosing, preventing or treating diseases associated with amyloidogenesis or amyloid-plaque formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antibody molecule; antibody; beta-A4 peptide; Abeta4; neuroprotective; nootropic; antiparkinsonian; gene therapy; amyloidogenesis; amyloid-plaque formation; beta-amyloid plaque; immunisation; dementia; allaheimer's disease; motor neuropathy; Down's syndrome; Creutzfeldt Jacob disease; hereditary cerebral haemorrhage; amyloidosis; Parkinson's disease; HIV-related dementia; amyotrophic lateral sclerosis;
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Pred. No. 1.7e-37;
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                                                                                                                                                                                                           Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cc environment comprising assessing the intracellular interaction between the immunoglobulin domain and the target by monitoring the signal. The method comprises providing a first molecule and a second molecule, where stable interaction of the first and second molecules leads to the companient of a signal; providing a single intracellular immunoglobulin domain which is associated with the first molecule, where the single immunoglobulin domain is free of complementary immunoglobulin domains; providing an intracellular target which is associated with the second molecule, such that association of the immunoglobulin domain and the carget leads to stable interaction of the first and second molecules and generation of the signal; and assessing the intracellular interaction between the immunoglobulin domain and the target by monitoring the signal. The methods are useful for determining the ability of an intracellular interaction of the signal. The methods are useful for determining the ability of an intracellular interaction of the signal. The present sequence is that of a single chain variable fragment (scFV) protein which was used to illustrate the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
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                                                                                                                                                                                                                                                                                                             ADQ75230 standard; protein; 119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention relates to a novel method of determining the ability of an immunoglobulin single domain to bind to a target in an intracellular
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Determining the ability of an immunoglobulin single domain to bind to target in an intracellular environment by assessing the intracellular interaction between the immunoglobulin domain and the target by monitoring the signal.
15-NOV-2002; 2002GB-00026731
                                  14-NOV-2003; 2003WO-GB004964
                                                                    03-JUN-2004.
                                                                                                      WO2004046189-A2
                                                                                                                                                                       immunoglobulin;
                                                                                                                                                                                                          Immunoglobulin heavy chain variable domain
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                                                                                                                                                                                                                                                                                                                                                                                                                    SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       QVQLVESGGGLVQPGGSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAFIRNDGSNEYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EVOLVESGGDFVQFGGSLRVSCAASGFAFSHYAMSWVRQAFGKGLEWVAYISSGGSGTYY
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                                                                                                                                                                         variable domain
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78.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 492; DB 8;
Pred. No. 1.2e-37;
8; Mismatches 18
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                                                                                                                                                                                                          A28 sequence
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RESULT 27
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Best Local S
Matches 93
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15-NOV-2002; 2002GB-00026723
15-NOV-2002; 2002GB-00026727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a method of preparing a double stranded nucleic acid, which encodes an immunoglobulin comprising providing a set of three or more overlapping oligonucleotides which anneal to form the + and - strands of a nucleic acid that encodes at least part of an immunoglobulin variable domain. The methods are useful for preparing nucleic acids, preferably immunoglobulin genes. This sequence represents one of the immunoglobulin variable heavy chain produced by the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Preparing a double stranded nucleic acid comprises providing a set o three or more overlapping oligonucleotides which anneal to form the strands of a nucleic acid that encodes at least part of an immunoglobulin variable domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antibody; variable chain; cytostatic; cytoplasmic degradation; intracellular relocation; specific antigen positive cancer; lelymphoma; intracellularly binding immunoglobulin; BCR-ABL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VH chain clone A28 of an intracellularly binding immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 119 AA;
                                                                                                               WPI; 2003-779088/73.
                                                                                                                                                                  Lobato-Caballero
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                                                                                                                                                                                                                        (MEDI-) MEDICAL RES
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78.2%;
                                                                                                                                                                  Rabbitts
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Pred. No. 1.2e-37;
8; Mismatches 18
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Use of an intracellularly binding immunoglobulin comprising at least antibody variable chain, in preparing a medicament for degrading one more specific antigens, or for treating specific antigen positive car

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               environment. Specifically, it refers to antibodies that can form an insoluble complex with a cognate antigen, such that it can then be target for degradation via the lysosome or proteosome systems. The present invention describes the specific target antigen as the oncognic fusion protein BCR-ABL or the RAS antigen, such that this method can be used to prepare a cytostatic medicament for the cytoplasmic degradation or intracellular relocation of such an antigen or for the treatment of the specific antigen positive cancer i.e. leukaemia or lymphoma. Furthermore, the immunoglobulins may also be used for therapeutic, prophlylactic or diagnostic applications both in vitro and in vivo, as well as for assay and reagent applications or in functional genomics. This polypeptide sequence is a variable heavy chain (VI) framework region of an
                                                                                                                                                                                                                                                                                                                                                                                 Human; novel protein; Antianaemic; osteopathic; antiinflammatory; immunomodulatory; cytostatic; neuroprotective; vulnerary; nootropic; anticonvulsant; antiarthritic; neuroprotective; antifungal; antiviral; antibacterial; antiallergic; dermatological; haemostatic; antialsthmatic; thrombolytic; immunogen; antibody; gene therapy; neurological disorder; Parkinson's disease; inflammatory disorder; cancer; asthma; osteoporosis
                                                        WPI; 2001-451939/48
N-PSDB; AAS22625.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 121 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to novel immunoglobulin molecules that comprise least one antibody variable chain VH or VL framework region and are capable of binding to a specific antigens within an intracellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              e.g. leukemia.
Isolated polypeptides useful for treating anti-inflammatory diseases, nervous system disorders, and for regenerating bone and cartilage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human novel protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24-OCT-2001
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                                                                                                                                                                                    25-JAN-2000; 2000US-00491404
                                                                                                                                                                                                                     25-JAN-2001; 2001WO-US002623.
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                                                                                                                                                (HYSE-)
                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119
                                                                                                                                                                                                                                                                                                                                                                regeneration; immune
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                                                                                                            Liu C,
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78.2%;
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Pred. No. 1.
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01-JUL-2002; 2002US-0392382P 26-JUN-2003; 2003WO-US019908 08-JAN-2004 WO2004003144-A2 Homo sapiens cancer; human. diabetes; non-insulin

dependent

diabetes; insulin

dependent diabetes;

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RESULT 29
AD158099
ID AD15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC equencing, for chromosome or gene mapping, in the production of crecombinant proteins, and in generating anti-sense DNA or RNA and in gene therapy. Polypeptides of the invention can be used to target drugs to a cumour, in assays to determine biological activity, to raise can tibodies/elicit an immune response, to determine quantitative protein clavels, as tissue markers, and to isolate receptors or ligands. CC entibodies, stem cell disorders, regenerating bone, cartilage, tendon, cligament and/or nerve tissue, wound healing, treating platelet cc disorders, stem cell disorders, regenerating bone, cartilage, tendon, cligament and/or nerve tissue, wound healing, treating burns, promoting ct the proliferation, differentiation and survival of stem cells, as a contraceptive, treating osteoporosis and osteoarthritis, anaemia, callarosis, stroke, immune deficiencies resulting from bacterial, viral or fungal infection or from autoimmunity, cancer, allergy, asthma, graft-creating osteomes, haemophilia, thrombosis, anti-inflammatory contenders, anti-of-the invention of invention of infection. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 313 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mammal and prevention of disorders caused by the aberrant protein expression or activity. The polypeptides can be used as molecular weight expression or activity. The polypeptides can be not been polypeptides markers, food supplements, and in antibody production. The polypeptides are used to identify compounds which bind to the polypeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 4; Page 630-631; 894pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   antibody; regeneration IV; Reg IV; single chain antibody fragment; scFv;
inflammatory bowel disorder; ulcerative colitis; Crohn's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reg IV-specific single chain antibody fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADI58099;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADI58099 standard; protein; 251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                represents a protein of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61
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74.8%;
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3.2e-37;
hes 15;
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Matches 94
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADJ95639 standard;
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N-PSDB; ADI58165.
Antibodies and antibody fragments recognising human insulin-like growth
                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
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                                                     WPI; 2003-854482/79
                                                                                                                                                                                                          30-APR-2002; 2002JP-00129046
                                                                                                                                                                                                                                                                                                                13-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                   WO2003093317-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               insulin-like growth factor; IGP-I;
                                                                                                                                                                                                                                                             30-APR-2003; 2003WO-JP005505
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                                                                                                      Nakamura
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Pred. No. 2.8e-37;
1; Mismatches 14
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                                                                                                                                                                                                                                                                                      Bardroff M,
Loehning C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antibody molecule; antibody; beta-A4 peptide; Abeta4; neuroprotective; nootropic; antiparkinsonian; gene therapy; amyloidogenesis; amyloid-plaque formation; beta-amyloid plaque; immunisation; dementia; Alzheimer's disease; motor neuropathy; Down's syndrome; Creutzfeldt Jacob disease; hereditary cerebral haemorrhage; amyloidosis; Parkinson's disease; HIV-related dementia; amyotrophic lateral sclerosis;
                                                                                                                             New antibody molecule capable of specifically recognizing two regions the beta-A4 peptide, useful for diagnosing, preventing or treating diseases associated with amyloidogenesis or amyloid-plaque formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      factor for treatment and diagnosis of cancer, diabetes and rheumatoid \mbox{arthritis.}
                                                   Example 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-FEB-2003; 2003WO-EP001759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003070760-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           neuronal disorder; aging.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anti-Abeta antibody related amino acid sequence SEQ ID NO:89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA89974;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADA89974 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                represents an insulin-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rheumatoid
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                                                                                                     (e.g. dementia).
                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-FEB-2002; 2002EP-00003844.
                                                                                                                                                                                                                                                                                                                                                                        (MORP-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention comprises antibodies and antibody fragments which bind to inhibit the function of human insulin-like growth factor (IGF-I /or IGF-II). The antibodies and antibody fragments of the invention useful for the treatment, prevention, and diagnosis of diabetes, umatoid arthritis and cancer. The present amino acid sequence resents an insulin-like growth factor-related protein of the
                                                                                                                                                                                                                                        2003-663848/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sapiens.
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95; Conser
                                                                                                                                                                                                                                                                                                                                                                        HOFFMANN LA ROCHE & CO
MORPHOSYS AG.
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                                                   Page
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                                                                                                                                                                                                                                                                                      Bohrmann B,
Loetscher H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IJ
                                                 53; 312pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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79.8%;
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                                                                                                                                                                                                                                                                                         Brockhaus M,
Nordstedt C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 126
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7:
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Pred.
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Mismatches
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No. 1.6e-37;
                                                                                                                                                                                                                                                                                                                  Huber W,
                                                                                                                                                                                                                                                                                            Rothe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15;
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describes an antibody molecule

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capable

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AAR95216
ID AAR95
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FT Domai
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC specifically recognising two regions of the beta-A4 peptide/Abeta4. The CF first region comprises the amino acid sequence Ala-Glu-Phe-Azg-His-Asp-CC ser-Gly-Tyr ADAB9886 or its fragment, and the second region comprises the CC amino acid sequence Val-His-His-Gln-Lys-Leu-Val-Phe-Phe-Ala-Glu-Asp-Val-CC gly ADA89887 or its fragment. Also described: (1) a nucleic acid molecule encoding (1); (2) a vector comprising the nucleic acid of (1); (3) a host CC culturing the host cell of (3) under conditions that allow synthesis of CC (1) and recovering (1) from the culture; (5) a composition comprising (1); (2) or vector of (2); (4) preparing (1), comprising (1); (2) or host cell of (3); (7) optimising (1); (2) or host cell of (3); (7) optimising (1); (8) testing the resulting Fab optimised clones; (10) expressing CC (1); (8) testing the resulting Fab optimised clones; (10) expressing CC of selected, optimised clones; (11) preparing a pharmaceutical composition, comprising optimisation of (1), and formulating the optimised antibody/antibody molecule with a carrier; and (12) a pharmaceutical composition of preparing a pharmaceutical composition in preparing a pharmaceutical composition for the used in gene therapy. The antibody molecule (1), nucleic acid molecule, vector or host is used in preparing a pharmaceutical composition for the disease mentioned above. The antibody is used for the disease mentioned above. The antibody is used for the disease is of the disease and/or amyloid plaque formation. The antibody molecule composition for the disease is disease, hereditary cerebral haemorrhage with camposition of the disease; hereditary cerebral haemorrhage with camposition of the present sequence is used in the exemplification of the present invention.
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Best Local (
                                                                         Key
Domain
                                                                                                                                                                    Antibody; fusion protein; single chain; inhibition; tumour; diagnosis; detection; imaging; immunotoxin; targetting; assay; immunoassay; Lewis(Y) carbohydrate antigen.
 Domain
                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                Human foetal immunoglobulin 56P1'CL variable heavy chain.
                                                                                                                                                                                                                                                                                                                             AAR95216;
                                                                                                                                                                                                                                                                                                                                                               AAR95216 standard; protein; 119
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                                     Domain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA;
                                                                                                                                                                                                                                                                                     (first entry)
                     31. .35
/label= CDR 1.
50. .66
50. .66
/label=
99. .108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                126
                                                                                           Location/Qualifiers
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                 CDR 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11; Mismatches
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ADA43059
ID ADA43
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Best Local S
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Fitzgerald I
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28-OCT-1994;
28-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A novel recombinant DNA molecule which encodes a single chain fusion protein or antibody comprising the Fv region of both the light and heavy chains of an antibody (Ab) fused together, and an effector molecule, where the fusion protein or Ab has the binding specificity of monoclonal Ab (MAb) B1, B3 or B5, can be used for the production of such fusion proteins or antibodies. The fusion proteins can be used in compositions as an immunotoxin to inhibit tumour cell growth. The single chain lewis(Y) carbohydrate antigen in a patient. The antibodies are also useful as multiple targetting moisties, providing at least 2 kinds of biological activity. They can also be used in diagnostic assays and for the imaging of tumours when attached to a radiolabel and for the pathological diagnosis of tumours. Humanised antibodies are less immunogenic than the mouse MAbs B1, B3 and B5, making them more suitable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Single chain fusion proteins and antibodies - useful to diagnose and treat cancer, specifically bind Lewis(Y) related carbohydrate antigen.
                                                                                      Homo sapiens
                                                                                                                                         myeloma; lymphoma;
                                                                                                                                                                       cytostatic; immunosuppressive; antiallergic; human leukocyte antigen; HLA-DR; antibody; hybridoma; diagnosis; cancer; allergy; sarcoma;
                                                                                                                                                                                                                                                             Human antibody HD4 heavy chain protein
                                                                                                                                                                                                                                                                                                                      20-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                    ADA43059 standard; protein; 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 119 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1996-251462/25.
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                             WO2003033538-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for long term treatment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARRSARTYYFDYWGQGTLVTVSS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 QVBLVESGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLEWVAVISYDGSNKYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 EVOLVESGGDFVOPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13; Fig 11A; 116pp;
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D, Brinkmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                      (first entry)
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94US-00331397.
94US-00331398.
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                                                                                                                                               leukaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        78.4%;
79.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Padlan
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Pai L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 489; DB 2;
Pred. No. 2.2e-37;
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Best Local Similarity
Matches 96; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      binding antibodies and their functional fragments and soluble domains. The specification also includes: a) hybridomas producing the antibodies; b) nucleic acids encoding all or part of the antibodies; c) expression vectors containing the nucleic acids; d) hosts transformed by the vectors; e) method for preparation of the antibodies by culture of the transformed hosts; and f) agents for the prevention, treatment and diagnosis of cancer and allergies, containing the antibodies. The antibodies are used in the prevention, treatment and diagnosis of allergies including breast, colon, kidney, stomach, ovary, pancreas, uterus, oesophagus, liver, skin, bladder and vascular cancer, pancreas, myeloma, lymphoma (including T-cell lymphoma, Hodgkin's lymphoma and non-Hodgkin's lymphoma) and leukaemia (including chronic and acute lymphocytic leukaemia). This sequence represents the heavy chain of the HD4 antibody targeted to the HLA-DR chains.
                                                                                                Anti-human CTLA-4 sFv; single chain antibody; phage; human CTLA-4; membrane-associated protein; ligand; activated T-cell; B7; CD28; co-stimulatory signal; T-cell proliferation; xenograft; organ transplant; xenograft-specific immunosuppression.
                                                                                                                                                                                                                  07-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to novel human leukocyte antigen DR (HLA-DR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      \ensuremath{\mathsf{HLA}}\xspace{-\mathsf{DR}} binding antibodies useful as immunosuppressant agents treatment and prevention of cancer and allergies.
                                                                                                                                                                                Anti-human CTLA-4 sFv.
                                                                                                                                                                                                                                                                               AAY15124 standard;
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                  Region
                                                                  Homo sapiens
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Pred. No. 2.7e-37;
9; Mismatches 14
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                       Anti-murine CTLA-4 sPv, M1 sFv, single chain antibody, murine CTLA-4 membrane-associated protein; chimeric construct; extracellular domain human CDB, ligand; activated T-cell; co-stimulatory signal; donor B7 recipient CD28; T-cell proliferation;
                                                       30-APR-1998;
                                                                                                           30-APR-1999;
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                                                                                                                                                                                                                                                                       Synthetic
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     (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000-038815/03.
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Pred. No. 4.6e-37;
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RESULT 36
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                                                                  Novel antibody, useful for treating, preventing or ameliorating inflammatory bowel disorder, cancer of the gastrointestinal tradiabetes (non-insulin dependent diabetes or insulin dependent diabetes)
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)B; ADI58124.
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                          NO 18;
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79.8%;
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                        324pp; English
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Pred. No. 4.6e
6; Mismatches
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.6e-37;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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Matches 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    useful for treating, preventing and ameliorating: inflammatory bowel disorders (e.g. ulcerative colitis or Crohn's disease), diabetes (e.g. non-insulin dependent diabetes or insulin dependent diabetes), and ca of the gastrointestinal tract. The antibody of the invention is also useful for detecting the expression of a Reg IV protein. The present amino acid sequence represents an scFv of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; antibody; tissue inhibitor of metalloprotease-1; TIMP-1; VLCDR3; matrix metalloprotease; MMP; variable heavy chain; VHCDR3; hepatotropic; variable light chain; cytostatic; nephrotropic; cardiant; liver fibrosis; alcoholic liver disease; cardiac fibrosis; acute coronary syndrome; lupus nephritis; glomerulosclerotic renal disease; lung cancer; lupus nephritis; glomerulosclerotic renal disease; lung cancer;
                                                                                                                                    New human anti-TIMP-1 (tissue inhibitor of metalloprotease-1) antibodies, for diagnosing or ameliorating the symptoms of a disorder in which TIMP-1 is elevated, e.g. liver fibrosis, benign prostate hypertrophy or lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 248 AA;
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                                                                                                                                                                                                                                                                                                                                             Pan C,
                                                                                                                                                                                                                                                                                                                                                                                                     (FARB )
(MORP-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human anti-TIMP-1 antibody heavy chain #17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          idiopathic pulmonary fibrosis; benign prostate hypertrophy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGT--YYFDSWGQGTLLTVS
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                                                                                                                                                                                                                                                                                                                                                Knorr AM, Schauer
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No. 4.8e-37;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC antibody, which binds to a tissue inhibitor of metalloprotease-1 (TIMP-1) CC and neutralises a matrix metalloprotease (MMP)-inhibiting activity of TMP CC -1. The antibody comprises a variable heavy chain (VHC)DR3 region and a CC variable light chain (VLC)DR3 region. An antibody preparation of the CC civity. The human antibody is useful for decreasing an MMP-inhibiting CC activity of a TIMP-1. It is especially useful for ameliorating the CC symptoms of a disorder in which TIMP-1 is elevated, e.g. liver fibrosis, CC alcoholic liver disease, cardiac fibrosis, acute coronary syndrome, lupus CC mephritis, glomeruloselerotic renal disease, idiopathic pulmonary CC fibrosis, benign prostate hypertrophy, lung cancer or colon cancer. The CC antibody is also useful for detecting a TIMP-1 in a test preparation, or in diagnosing a disorder in which a TIMP-1 level is elevated. The CC sequences shown in ABRO1502-ABRO1545 represent the heavy chain regions of CC a human anti-TIMP-1 antibody of the invention
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Best Local S
Matches 95
                 preparing a eclampsia. invention.
                                                    B. An antibody of the invention has hypotensive, and gynaecological
activity, and may have a use in gene therapy. The antibody is useful for
preparing a composition for treating or preventing hypertension or pre-
                                                                                                                                                  Claim
                                                                                                                                                                                   New antibody that specifically binds neurokinin B, useful for preparing composition for treating or preventing hypertension or preeclampsia.
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                                                                                                                                                                                                                                                                                                                                                                            30-MAY-2002; 2002US-0383802P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hypertension;
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                                                                                                             The invention relates
                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                             2004-053456/05
                                                                                                                                                2; SEQ ID NO 32; 127pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               neurokinin B; hypotensive; gynaecological;
ion; pre-eclampsia; NKB.
                                                                                                                                                                                                                                                                                                  Ruben SM
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79.8%;
                                                                                                                                                                                                                                                                                                                                       SCI INC
                                                                                                             to a novel antibody specifically binding
                                   sequence
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Pred. No. 4.7e-37;
9; Mismatches 12
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                    Humanised anti-L-selectin antibody HuDreg 55 comprises 2 heavy chains each having the sequence given in AAW13564 and 2 light chains each having the sequence given in AAW13563. These are encoded by the cDNA clones given in AAR61281 and AAR61280. HuDreg 55 can be used to prevent multiple organ failure associated with polytrauma and for the prevention of acute organ damage associated with extracorporeal blood circulation. The antibody inhibite interaction between the carbohydrate-recognising domain of the selectin and the corresponding cell surface receptor. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                 17-AUG-1995;
19-SEP-1995;
27-DEC-1995;
                                                                                                                                                                  Using anti-selectin antibody to prevent acute organ damage and multiple organ failure - during extracorporeal circulation or following polytrauma, e. g. haemorrhagic-traumatic shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 248
                                                                                                                                           Disclosure; Page 34-36; 52pp; English.
                                                                                                                                                                                                                                                     Haselbeck A,
                                                                                                                                                                                                                                                                               (PROT-)
                                                                                                                                                                                                                                                                                                                                                                 14-AUG-1996;
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                                                                                                                                                                                                                                                                              PROTEIN DESIGN LABS INC. BOEHRINGER MANNHEIM GMBH.
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95EP-00114696.
95US-00578953.
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Best Local
The present invention describes a composition (1), comprising a polypeptide comprising an antibody-based antigen-binding domain of human composition with binding specificity for an antigen expressed on the surface of a cell, where treating cells expressing the antigen with the polypeptides leads to suppression of an immune response, and the IC50 for the suppression of immune response is 1 microM or less. (1) has antirheumatic, antiarthritic, neuroprotective, antiinflammatory, antiarthritic, immunosuppressive, dermatological, antithyroid, nephrotropic, thyromimetic and hepatotropic activities, and can be used as suppressor of immune response. (1) is useful for suppressing IL-2 secretion by a cell, the interaction of a cell of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunomodulatory human MHC class II antigen-binding protein; HLA; human leukocyte antigen; immune system; immunosuppression; antibody; major histocompatibility complex; antirheumatic; antipsoriatic; neuroprotective; antiinflammatory; antidiabetic; antipsoriatic; immunosuppressive; dermatological; antithyroid, nephrotropic; psoriasis; thyromimetic; hepatotropic; immune response suppressor; narcolepsy; rheumatoid arthritis; juvenile arthritis; multiple sclerosis; insulitis; grave's disease; insulin-dependent diabetes; Hashimoto's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Composition for suppressing immune response, treating diseases of immune system, has polypeptide comprising antibody-based antigen-binding domain of human composition, which binds antigen expressed on a cell surface.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 systemic lupus erythematosus; ankylosing spondylitis; myasthenia gravis; transplant rejection; graft versus host disease; pemphigus vulgaris; glomerulonephritis; thyroiditis; pancreatitis; primary biliary cirrhosis; irritable bowel disease; Sjogren's syndrome.
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06-OCT-2000; 2000US-0238762P.
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ilarity 82.4%;
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Pred. No. 1.2e-36;
6; Mismatches 12
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Matches 95
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ABA92469 to ABA92474 and ABB57457 to ABB57590 represent sequence used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immune system with another cell, immunosuppressing a patient and for killing a cell expressing an antigen, human leukocyte antigen (HLA)-DR the surface of the cell, where neither cytotoxic entities nor immunological mechanisms are needed to cause or lead to the killing. (1 (optionally linked to cytotoxic or immunogenic agent) is useful for preparing a pharmaceutical preparation for the treatment of rheumatoid arthritis inventie architis more treatment of recommendations.
                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                        exemplification of the present invention
                                      119
121 S 121
                                                                               61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARSSMPMYGEGFDLWGQGTLVTVS
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78.5%;
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Pred. No. 3.4e-37;
7; Mismatches 17;
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Search completed: December 17, 2004, 18:29:09 Job time : 363.68 secs

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Result
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Maximum
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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109.302 Million cell updates/sec
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     GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
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US-09-764-304-8
US-09-764-31-33C-4
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US-08-331-398A-46
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US-08-331-398A-20
US-08-755-804A-26
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US-08-755-378A-20
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US-09-705-686-21
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Sequence 31, Appli
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## ALIGNMENTS

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RESULT 2
US-09-764-304-18
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US-09-225-322B-18
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PRIOR FILING DATE: 1995-05-31
PRIOR PELICATION NUMBER: US 08/408,133
PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US07/947,674
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Best Local Simi
Matches 109;
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Patent No. 6437098
Patent No. 6437098
GENERAL INFORMATION:
APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIXAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/225,322B
CURRENT FILING DATE: 1999-01-05
                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 1991-09-18

NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 18
LENGTH: 130
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:cDNA KM-641
                                                                                                                                                                                                                                                                                         Local Similarity
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                                                                                                      SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119
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91.6%;
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Pred. No. 1e-49;
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INFORMATION:

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RESULT 3
US-09-225-322B-8
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; OTHER INFORMATION: cDNA KM-641
US-09-764-304-18
                                                                                                                                                                                                                    FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/09/225,322B
CURRENT FILING DATE: 1999-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1994-08-17
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EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER FILLING DATE: 1995-05-31
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER FILLING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/292,178
EARLIER FILLING DATE: 1994-08-17
EARLIER APPLICATION NUMBER: US07/947,674
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LENGTH: 130
TYPE: PRT
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Best Local Similarity
SOFTWARE:
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CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 2001-01-19
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APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEAMA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
                                                                                          PRIOR APPLICATION NUMBER: US07/947,674
PRIOR FILING DATE: 1992-09-17
PRIOR APPLICATION NUMBER: JP 3-238375
PRIOR FILING DATE: 1991-09-18
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APPLICANT: HASEGAWA, MAMORU
                                                              NUMBER OF SEQ ID NOS: 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    71
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                               PatentIn Ver. 2.0
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91.6%;
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Pred. No. 1e-49;
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RESULT 5 US-09-648-067A-15

Sequence 15, Application US/09648067A Patent No. 6627196

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; LENGTH: 130
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; COTHER INFORMATION: Description of Artificial Sequence: cDNA
US-09-225-322B-8
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Best Local S
Matches 108
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EARLIER FILING DATE: 1992-09-17
EARLIER APPLICATION NUMBER: JP 3-238375
EARLIER FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 8
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Matches
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Patent No. 6495666
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EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER FILING DATE: 1995-05-31
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/292,178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
FILE REFERENCE: 249-101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EARLIER FILING DATE: 1994-08-17
                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                  OTHER INFORMATION: cDNA KM-641
                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                               LENGTH: 130
                                                                                                                                                                       Local
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                    61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119
                                                                          11 EVTLVESGGDFVKPGGSLKVSCAASGFAFSHYAMSWVRQTPAKRLEWVAGISSGGSGTYY 70
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71 SDSVKGRFTISRDNAKNTLYLQMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS 129
                                                                                                1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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Pred. No. 9.6e-49;
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Pred. No. 9.6e-49;
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RESULT 6
US-07-934-373C-4
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Best Local Similarity
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SEQ ID NO 15
LENGTH: 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4, Application US/07934373C Patent No. 5821337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/09/648,067A
CURRENT FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/151,018
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: US 60/213,822
PRIOR FILING DATE: 2000-06-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Paul J. Carter
APPLICANT: Leonard G. Presta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
FILE REFERENCE: P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Baughman, S
APPLICANT: Shak Steven
TELEFAX: 650/952-9881
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                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
COMPUTER: OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/
FILING DATE: 15-UN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/715272
FILING DATE: 14-UN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 1 SOUTH San Fra
                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wandy M
                                                                                                                                                                                                                                                                                                                                          SOFTWARE: WinPatin (Ge
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: IN NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                            TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                 COMPUTER: 18" .. PC-DOS/NO ... OPERATING SYSTEM: PC-DOS/NO ... OPERATING SYSTEM: PC-DOS/NO ... PC-DOS/NO DATA:
                                                                              NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: PO
                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/07/934,373C FILING DATE: 21-Aug-1992 CLASSIFICATION: 530
                                         TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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San Francisco
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                                                                                 P0709P2
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Pred. No. 8.3e-42;
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; TYPE: Am:
; TOPOLOGY:
US-07-934-373C-4
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Best Local Similarity
Matches 97; Conserv
 Query Match
Best Local Similarity
Matches 97; Conserv
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APPLICANT: Paul J. Carter
APPLICANT: Leonard G. Presta
                                                                                                                            INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: PO
                                                                                                                                                                                                                                          FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
                                                                                                                                                                                                                                                                                                      FILING DATE: 15-JUN-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/05126
                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                                                                              TYPE: Amino Acid
                                                                                                                                                                                                                        REGISTRATION NUMBER: 40,378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/437,642B FILING DATE: 09-May-1995 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ITLE OF INVENTION: Immunoglobulin Variants
                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
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                                                                                                                                                                           TELEPHONE:
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Amino Acid
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milarity 80.8%;
Conservative
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 Score 495.5; DB
Pred. No. 1.5e-41
7; Mismatches 1
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Pred. No. 1.5e-41;
                                 DB 3;
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                               Length 120;
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RESULT 9
US-09-705-686-4
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Sequence 4, Application US/09705686
Patent No. 6639055
GENERAL INFORMATION:
APPLICANT: Carter, Paul J.
Presta, Leonard G.
TITLE OF INVENTION: Method for Making Humanized Antibodies
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 4:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: PO-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WinPatin (Ge
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/715272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Presta, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/146,206C FILING DATE: 17-No. 6407213-1993 CLASSIFICATION: 530
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                                                                                                                                                                                                                                                                                                                                                   Local
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                                                                                                                                                                                           13
                                                                                                                                                                                                           61 SDSVKGRFTISRDNSKNTLYLØMRSLRAEDSAVYFCTRVKLGTY-YFDSWGQGTLLTVSS 119
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80.8%;
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Method
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for Making Humanized Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P0709P1
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US-09-705-686-4
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Best Local Similarity 80.8
Matches 97; Conservative
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: 650/225-1994
TELEPAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 120 amino acids
TYPE: Amino Acid
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
APPLICATION NUMBER: 07/715272
FILING DATE: 14-UN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: 90709PID3
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MEDIUM TYPE: 3.5 inch, 1.44 Mb flops

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WinPatin (Genentech)

CURRENT APPLICATION NUMBER: US/09/705,686

FILING DATE: 02-No. 6639055-2000

CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/705,392A
FILING DATE: 02-No. 6719971-2002
                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Carter, Paul J.
Presta, Leonard G.
TITLE OF INVENTION: Method for Making Humanized Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Genente
                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Genente
                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Carter,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 SDSVKGRFTISRDNSKNTLYLØMRSLRAEDSAVYFCTRVKLGTY-YFDSWGQGTLLTVSS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 ADSVKGRFTISRDDSKNTLYLOMNSLRAEDTAVYYCARDRGGAVSYFDVWGQGTLVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 EVOLVESGGGLVOPGGSLRLSCAASGFTFSDYAMSWVRQAPGKGLEWVAVISENGSDTYY
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STATE: California
                                                                                                                                                                                                                          CITY: South San Fi
STATE: California
                                                                                                                                                                              ZIP: 94080
                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                      STREET: 1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 1 DNA Way
                                                                                                                                                                                                     USA
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                                                                                                                                                                                                                                                                                           Genentech, Inc.
                                                                                                                                                                                                                                                 Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            inch, 1.44 Mb floppy disk
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Pred. No. 1.5e-41;
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ZIP: 94105-1492
ZIP: 94105-1492
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PacentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/331,398A
FILING DATE: 28-OCT-1994
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US 07/767,331
FILING DATE: 30-SEP-1991
PRIOR APPLICATION NUMBER: US 07/767,331
FILING DATE: 30-SEP-1991
PRIOR APPLICATION NUMBER: US 07/596,289
FILING DATE: 12-OCT-1990
FILING DATE: 12-OCT-1990
FILING DATE: 12-OCT-1990
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US-08-331-398A-46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 46, Application US/08331398A
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
TELEPAX: 650/922-9881
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Single Chain B3 Antibody Fusion Proteins TITLE OF INVENTION: and Their Uses (as amended) NUMBER OF SEQUENCES: 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
REFERENCE/DOCKET NUMBER: 015280-126110US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 07/715272
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE DESCRIPTION: SEQ ID NO: 4:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADSVKGRFTISRDDSKNTLYLOMNSLRAEDTAVYYCARDRGGAVSYFDVWGQGTLVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: Amino Acid
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/146206 FILING DATE: 17-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                               California
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One Market Plaza, Steuart Street Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Willingham, Mark
FitzGerald, David
Brinkmann, Ulrich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         79.4%;
80.8%;
                  38,498
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Pred. No. 1.5e-41;
7; Mismatches 15
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FILING DATE: 12-OCT-1990
ATTORNEY/AGENT INFORMATION:
NAME: HUNTEY, Tom
REGISTRATION UNMBER: 38,498
REFERENCE/DOCKET NUMBER: 015280-126120US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
TELEPAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
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TELEPHONE: (415) 543-960
TELEPAX: (415) 543-9043
INFORMATION FOR SEQ ID NO: 46:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Pastan
APPLICANT: Benhar
                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/331,3978
FILING DATE: 28-OCT-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/767,331
FILING DATE: 30-SEP-1991
                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: 1..119
OTHER INFORMATION: /note= "Human fetal immunoglobulin OTHER INFORMATION: 56P1'CL Variable Heavy chain (V-H)"
                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 EVOLVESGGDFVQFGGSLRVSCAASGFAFSHYAMSWVRQAFGKGLEWVAYISSGGSGTYY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 QVELVESGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLEWVAVISYDGSNKYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94105-1492
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5981726
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79.0%;
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Specific
Thereof
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                                                                                                                                                                                                       US 07/596,289
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 489; DB 1;
Pred. No. 6.4e-41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and Mutationally Stabilized Tumor Antibody Fragments, Fusion Protein
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RESULT 13
US-08-759-804A-46
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Best Local !
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION: APPLICANT: Pastan
                                      TELEPHONE: (415) 576-030 TELEFAX: (415) 576-030 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/759,804A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANI: APPLICANI: Pai, Lee
APPLICANT: Pai, Lee
TITLE OF INVENTION: Tumor-Specific Antibody ****
TITLE OF INVENTION: Fusion Proteins, and Uses Thereof
TITLE OF SECULENCES: 68
SEQUENCE CHARACTERISTICS:
LENGTH: 119 amino acid
                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen L.
REGISTRATION NUMBER: 32,7
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/596,289
FILING DATE: 12-OCT-1990
                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 536
PRIOR APPLICATION UNATA:
APPLICATION UNMBER: US 08/331,398
FILING DATE: 28-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein
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                                                                                             TELECOMMUNICATION INFORMATION:
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OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                                                                                                                                                                                      FILING DATE: 30-SEP-1991
                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP:
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FitzGerald, David
Brinkmann, Ulrich
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                                                        576-0300
                                                                            576-0200
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79.0%;
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56P1'CL Variable Heavy chain (V-H)"
                                                                                                                                                                                                                                                                             US 07/767,331
                                                                                                                                     32,762
                                          46:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 489; DB 2;
Pred. No. 6.4e-41;
7; Mismatches 18
                                                                                                                   015280-126140US
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Patent No.
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Best Local Similarity
                                                          TELEFAX: (415) 543-504
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                          APPLICATION NUMBER: US 07/707,500
FILING DATE: 30-SEP-1991
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/596,289
                                                                                                                                                     APPLICATION NUMBER: US 07, FILING DATE: 12-OCT-1990
ATTORNEY/AGENT INFORMATION: NAME: Waber, Ellen Lauver REGISTRATION NUMBER: 32,71
                                                                                                                   REFERENCE/DOCKET NUMBER: 15
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
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LOCATION: 1..119
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: San Francisco
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TOPOLOGY: 11
              STRANDEDNESS:
                           LENGTH: 119 amino acids
TYPE: amino acid
                                                                                                       TELEPHONE:
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BENHAR,
PADLAN,
                                                                                       (415) 543-96
(415) 543-5043
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linear
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                                                                                                       543-9600
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79.0%;
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56P1'CL Variab
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Pred. No. 6.4e-41;
7; Mismatches 18
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ble Heavy chain (V-H)"
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 Query Match
                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION UMBER: PCT/US
APPLICATION NUMBER: PCT/US
FILING DATE: 29-OCT-193
ATTORNEY/AGENT INFORMATION:
NAME: Howells, Stacy L.
REGISTRATION UMBER: 34,84
REFERENCE/DOCKET NUMBER: F
                                                                                                                                                                                                                               TELEFAX: (619) 455-5110 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 90067
COMPUTER READABLE FORM:
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FEATURE:
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                                                                                                                 MMEDIATE SOURCE:
                                                                                                                                                                                                                                                                               ELECOMMUNICATION INFORMATION:
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/428,197 FILING DATE:
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OTHER INFORMATION: /note= "Human fetal immunoglobulin
OTHER INFORMATION: 56Pl'CL VH region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
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                                                                 NAME/KEY:
                                                                                                   CLONE: 18/2
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                                                 LOCATION:
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: California
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                                                                                                                                                                               amino acid
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1880 Century Park East - Suite 500
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                                                Peptide
1..125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Protein
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METHOD FOR STIMULATING PRODUCTION OF

VARIABLE REGION GENE FAMILY RESTRICTED ANTIBODIES THROUGH

VACCINATION WITH A B-CELL SUPERANTIGEN AND CONJUGATES

THEREOF
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79.0%;
 78.4%;
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Pred. No. 6.4e-41;
7; Mismatches 18
 Score 489;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Version #1.25
DB 2;
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Length 125;
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                                                                                                             Matches
                                                                                                                          Query Match
Best Local &
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                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (619) 455-510
TELEFAX: (619) 455-5110
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                        IMMEDIATE SOURCE:
CLONE: 18/2
                                                                                                                                                                                                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: FD
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Howells, Stacy L.
REGISTRATION NUMBER: 34,842
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TITLE OF INVENTION:
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LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                    ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCTKGQVLYYGSGSYHWFDPWGQGTL
61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKL-----GTY-YFDSWGQGTL 114
                                                                                                             95;
                                                                   1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                   amino acid
                                                 EVQLLESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYY
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1880 Century Park East - Suite 500
                                                                                                                                                                                                                                                                                                                                  125 amino acids
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                                                                                                             Conservative
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JENTION: METHOD FOR STIMULATING PRODUCTION OF
JENTION: VARIABLE REGION GENE FAMILY RESTRICTED
JENTION: VACCINATION WITH A B-CELL SUPERANTIGEN
                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                    peptide
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                                                                                                                                                                                                                                                                                                   single
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                                                                                                            78.4%; Sc.
76.0%; Pr
tive 12;
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                                                                                                            Score 489; DB 5;
Pred. No. 6.7e-41;
12; Mismatches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                FD-2630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred. No. 6.7e-41;
2; Mismatches 12
                                                                                                                                         Length 125
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AND CONJUGATES
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US-08-579-378A-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE: protein US-08-579-378A-20
                                                                                                                                                                                                                                                                                         Matches
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Best Local :
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APPLICATION NUMBER: EP 95112895.8

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: EP 95114696.8

FILING DATE: 19-SEP-1995

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 415-326-2422
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PATENTIN Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/579,378A
FILING DATE: 27-DEC-1995
CLASSIFICATION 424
PRIOR APPLICATION DATA: 08/160,074
APPLICATION NUMBER: US 08/160,074
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Co, Ma
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TITLE OF INVENTION: Huma
TITLE OF INVENTION: L-Se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Liebescheutz, Joe O.
REGISTRATION NUMBER: 37,505
REFERENCE/DOCKET NUMBER: 11823-002220
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 30-NOV-
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
FILING DATE: 01-DEC-
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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                                                                                                                                       20 EVQLVESGGGLVQPGGSLRLSCAASGFTFSTYAMSWVRQAPGKGLEWVASISTGGS-TYY
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79 PDSVKGRFTISRDNAKNTLYLOMNSLRAEDTAVYYCARDYDG--YFDYWGQGTLVTVSS 135
                                     61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119
                                                                                                                                                                                                                                                                                             98;
                                                                                                                                                                                    1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY 60
                                                                                                                                                                                                                                                                                                                               Similarity
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nilarity 82.4%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   135 amino acids
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415-326-2422
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                                                                                                                                                                                                                                                                                     Score 487.5; DB 3;
Pred. No. 1e-40;
6; Mismatches 12;
                                                                                                                                                                                                                                                                                                                                                         DB 3;
                                                                                                                                                                                                                                                                                                                                                             Length 135;
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RESULT 18

Sequence 21, Application US/07934373C Patent No. 5821337

GENERAL INFORMATION:

APPLICANT: Paul J. Carter APPLICANT: Leonard G. Presta

TITLE OF INVENTION: Immunoglobulin Variants

NUMBER OF SEQUENCES: 48 CORRESPONDENCE ADDRESS:

ADDRESSEE: STREET: 1

1 DNA Way

Genentech, Inc

South San Francisco California

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US-07-934-373C-21
                     RESULT 19
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                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
PRIOR APPLICATION NUMBER: 08/578,953
APPLICATION NUMBER: 08/578,953
FILING DATE: 27-Dec-95
APPLICATION NUMBER: EP 95 112 895.8
FILING DATE: 17-Aug-95
APPLICATION NUMBER: EP 95 114 969.9
FILING DATE: 19-Sep-95
ATTORNEY/AGENT INFORMATION:
NAME: NOTMON D. HANBOR
REGISTRATION NUMBER: 30,946
REGISTRATION NUMBER: BOER 1059-PCT
                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (212) 838-3884
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Computer Disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Martin, Ulrich, et al. TITLE OF INVENTION: Anti-selectin NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: ASCII
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: dou
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 805 Third Avenue CITY: New York STATE: New York COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: PCT/US96/13152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                          LENGTH:
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ADDRESSEE: Attn: Norman D. Hanson
                                                                          60
                                                                                              61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119
                                                                                                                                                                     1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                                                                  1 EVQLVESGGGLVQPGGSLRLSCAASGFTFSTYAMSWVRQAPGKGLEWVASISTGGS-TYY 59
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                                                                          PDSVKGRPTISRDNAKNTLYLOMNSLRAEDTAVYYCARDYDG--YFDYWGOGTLVTVSS 116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (212) 688-9200
                                                                                                                                                                                                                                                                                                                                                       double
                                                                                                                                                                                                                                          78.1%;
82.4%;
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                                                                                                                                                                                                                           6; Mismatches
                                                                                                                                                                                                                                          Pred. No. 3.9e-40;
                                                                                                                                                                                                                                                             DB 5;
                                                                                                                                                                                                                            12;
                                                                                                                                                                                                                            Indels
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COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inc

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USA

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RESULT 20
US-08-437-642B-21
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                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Paul J. Carter
APPLICANT: Leonard G. Presta
TITLE OF INVENTION: Immunoglob
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21, Application US/08437642B Patent No. 6054297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/07/934,373C FILING DATE: 21-Aug-1992 CLASSIFICATION: 530 PRIOR APPLICATION DATA: APPLICATION NUMBER: PCT/US92/05126 FILING DATE: 15-JUN-1992 PRIOR APPLICATION DATA: APPLICATION NUMBER: 07/715272 APPLICATION NUMBER: 07/715272
                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
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LENGTH: 122 amino acids
TYPE: Amino Acid
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REFERENCE/DOCKET NUMBER: PO
TELECOMMUNICATION INFORMATION:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/437,642B
FILING DATE: 09-May-1995
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                              ADDRESSEE: Genentech, Inc
STREET: 1 DNA Way
CITY: South San Francisco
                                                                                                                                                                                      COUNTRY: U
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al Similarity 77.0%;
94; Conservative 1
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                                                                                                                                                                                                                                                                                                                                    Immunoglobulin Variants
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TOPOLOGY:
US-08-437-642B-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 6407213
GENERAL INFORMATION:
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Best Local Similarity
APPLICATION NUMBER: US/08/146,206C
FILING DATE: 17-No. 6407213-1993
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/715272
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 122 amino acid
                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                    APPLICANT: Carter, Paul J.
APPLICANT: Presta, Leonard G.
APPLICANT: Presta, Leonard G.
ATITLE OF INVENTION: Method for Making Humanized Antibodies
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: PO
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
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FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 15-JUN-1992 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 17-NOV-
PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
FILING DATE: 21-AUG-
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                                                                                                                                                                                                                                                                           CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                       ZIP: 94080
                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                 STREET:
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Amino Acid
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RESULT 22
US-09-705-686-21
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US-09-705-686-21
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                          INFORMATION FOR SEQ ID NO: 21: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: PO
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/705,686
FILING DATE: 02-No. 6639055-2000
CLASSIFICATION: <URKnown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
APPLICATION WIMBER: 07/715272
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 122 amino acids
TYPE: Amino Acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Carter, Paul J.
Presta, Leonard G.
TITLE OF INVENTION: Method for Making Humanized Antibodies
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
                SEQUENCE DESCRIPTION: SEQ ID NO: 21:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                         REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: P0709P1D3
                                                    LENGTH: 122 amino acids
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 94080
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CITY: South San Francisco
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77.0%;
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Pred. No. 1.4e-40;
0; Mismatches 15;
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US-09-705-392A-21
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APPLICATION NUMBER: 07/715272

FILING DATE: 14-JUN-1991

ATTORNEY/AGENT INFORMATION:

NAME: Lee, Wendy M.

REGISTRATION NUMBER: 40,378

REFERENCE/DOCKET NUMBER: P0709P1D1 REVISED

TELECOMMUNICATION INFORMATION:

TELEPHONE: 650/225-1994

TELEPHONE: 650/25-1994

TELEPAX: 650/952-9881

INFORMATION FOR SEQ ID NO: 21:

SEQUENCE CHARACTERISTICS:
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Best Local S
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Best Local Similarity
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MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/705,392A
FILING DATE: 02-No. 6719971-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Carter, Paul J.
Presta, Leonard G.
TITLE OF INVENTION: Method for Making Humanized Antibodies
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                                                                                                                                                        Similarity
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                                                            EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSVISGDGGSTYY 60
                                                                                                                                                                                                                                                                 TYPE: Amino Acid
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CITY: South San Francisco
                                                                                                                                                                                                                                                      TOPOLOGY: Linear
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                                                                                                                                                      77.8%; Score 485.5; DB 4
77.0%; Pred. No. 1.4e-40;
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                                                                                                                                                                          DB 4;
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RESULT 25
US-09-069-821-3
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Best Local S
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/
FILING DATE: 15-JUN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/934373
FILING DATE: 21-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
CLASSIFICATION DATA:
PRIOR APPLICATION UNMBER: 07/715272
FILING DATE: '14-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOTTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US9
FILING DATE: 19930820
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MEDIUM TYPE: 5.25 inch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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ADDRESSEE: Genentech,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Immunoglobulin Variants NUMBER OF SEQUENCES: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
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                                                                121 SS 122
                                                                                                118 SS 119
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                                                                                                                                               61 SDSVKGRFTISRDNSKWTLYLQMRSLRAEDSAVYFCTRVKLG---TYYFDSWGQGTLLTV
                                                                                                                                                                                                                                                           94;
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                                                                                                                              61 ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARGRVGYSLSGLYDYWGQGTLVTV
                                                                                                                                                                                                                      1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
                                                                                                                                                                                             1 EVOLVESGGGLVOPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSVISGDGGSTYY
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94; Conservative
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South San Francisco
California
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                                                                                                                                                                                                                                                                                                                                       linear
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SYSTEM: PC-DOS/MS-DOS
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77.0%;
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Pred. No. 1.4e-40;
0; Mismatches 15
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Patent No. 6323322
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 60/044,449
FILTING DATE: 30-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 30-APR-1998
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/067,341
                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOPTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/069,821
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: SINGLE-CHAIN ANTIGEN-BINDING PROTEINS TITLE OF INVENTION: CAPABLE OF GLYCOSYLATION, PRODUCTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: Bil
TOPOLOGY: not re
                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: KIM, JUDITH U.
REGISTRATION NUMBER: 40,679
REFERENCE/DOCKET NUMBER: 0977.2280003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U
ZIP: 20005
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 02-DEC-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: STERNE, KESSLER, STREET: 1100 NEW YORK AVE.,
250
                              110 GOGTLLTVSS 119
                                                                 190
                                                                                                                                   130 EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSVISGKTDGGST 189
                                                                                             59 YYSDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGT------YYFDSW 109
                                                                                                                                                          1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYIS--SGGSGT
                                                                                                                                                                                                      96;
GQGTLVTVSS
                                                                 YYADSVKGRFTISRDNSKNTLYLOMNSLRAEDTÄVYYCARGRXGXSLSGXYYYYHYFDYW
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WHITLOW, MARC
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not relevant
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                                                                                                                                                                                                                                                                                                                                                                                                                              (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-OCT-1997
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73.8%;
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                                                                                                                                                                                                    Score 485.5; DB 3
Pred. No. 3.4e-40;
9; Mismatches 14
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NW, SUITE 600
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RESULT 26

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RESULT 27
US-09-956-087-3
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION: (202)371-2600
TELEPAX: (202)371-2540
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/069,821

FILING DATE: <Unknown:
APPLICATION NUMBER: US 60/063,074

FILING DATE: 27-OCT-1997

APPLICATION NUMBER: US 60/050,472

FILING DATE: 23-JUN-1997

APPLICATION NUMBER: US 60/044,449

FILING DATE: 30-APR-1997

APPLICATION NUMBER: US 60/044,449

FILING DATE: 30-APR-1997

ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION UMBER: US/09/956,086

FILING DATE: 20-Sep-2001

CLASSIFICATION: <UNknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 263 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: No. 6743896 Relevant
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.
STREET: 1100 NEW YORK AVE., NW, SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: SINGLE-CHAIN ANTIGEN-BINDING PROTEINS

CAPABLE OF GLYCOSYLATION, PRODUCTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: FILPULA, DAVID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 33
                                                                                                                                110 GOGTLLTVSS 119
                                                                                                                                                                            190 YYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGRXGXSLSGXYYYYHYFDYW 249
                                                                                                                                                                                                                                                                130 EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSVISGKTDGGST 189
                                                                                                                                                                                                                                                                                                                                                        96;
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                                                                                       GQGTLVTVSS 259
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COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: KIM, JUDITH U. REGISTRATION NUMBER: 40,679
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                                                                                                                                                                                                                                                                                                                                                        Conservative
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73.8%;
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Pred. No. 3.4e-40;
9; Mismatches 14;
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Query Match
Best Local Similarity
Watches 96; Conserva
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; SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-956-087-3
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Patent No. 6743908
GENERAL INFORMATION:
APPLICANT: FILPULA, DAVID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/956,087

FILING DATE: 20-Sep-2001

CLASSIFICATION NUMBER: US/09/956,087

FILING DATE: 20-Sep-2001

PRIOR APPLICATION CURRENCE

APPLICATION UNMER: 09/069,821

FILING DATE: 1998-04-30

APPLICATION NUMBER: US 60/063,074

FILING DATE: 27-OCT-1997

APPLICATION NUMBER: US 60/050,472

FILING DATE: 23-UN-1997

APPLICATION NUMBER: US 60/044,449

FILING DATE: 33-JUN-1997

APPLICATION NUMBER: US 60/047,449

FILING DATE: 33-JUN-1997

APPLICATION NUMBER: 40,679

REFERENCE/DOCKET NUMBER: 0977.2280003

TELECOMMUNICATION INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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  250
                                                                                                                                                                                                 130 EVQLVESGGGLVQPGGSLKLSCAASGFTFSSYAMSWVRQAPGKGLEWVSVISGKTDGGST 189
                                                                                                190 YYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGRXGXSLSGXYYYYHYFDYW
                                                                                                                               59 YYSDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGT------YYFDSW 109
                                                                                                                                                                                                                                1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYIS--SGGSGT
                                                 GQGTLLTVSS 119
GQGTLVTVSS 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C. STREET: 1100 NEW YORK AVE., NW, SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LEE, LIHSYNG S.
OF INVENTION: SINGLE-CHAIN ANTIGEN-BINDING PROTEINS
CAPABLE OF GLYCOSYLATION, PRODUCTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 263 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (202)371-2600
                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WHITLOW, MARC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SHORR, ROBERT
                                                                                                                                                                                                                                                                                                                            77.8%;
73.8%;
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Pred. No. 3.4e-40;
9; Mismatches 14
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                                                                                                        249
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RESULT 28 US-09-420-592A-6

Sequence 6, Application US/09420592A

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                                                                                    SEQ ID NO 6
LENGTH: 283
TYPE: PRT
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SEQ ID NO 6
LENGTH: 283
TYPE: PRT
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APPLICANT: Whitlow, Marc D.
APPLICANT: Whitlow, Marc D.
APPLICANT: Whitlow, Marc D.
APPLICANTON: No. 633396el Method for Targeted Delivery of Nucleic Acids
FILE REFERENCE: 0977.2300001
CURRENT APPLICATION NUMBER: US/09/420,592A
CURRENT FILING DATE: 1998-10-19
PRIOR APPLICATION NUMBER: 60/104,949
PRIOR FILING DATE: 1998-10-20
NUMBER OF SEQ ID NOS: 13
                                                                                                                                                                                                                                                                                                                      APPLICANT: Filpula, David R.
APPLICANT: Wang, Macliang
APPLICANT: Whitlow, Marc D.
TITLE OF INVENTION: No. 6692942el Method for Targeted Delivery of Nucleic Acids
FILE REFERENCE: 0977.2300003
                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/985,442
CURRENT FILING DATE: 2001-11-02
PRIOR APPLICATION NUMBER: 09/420,592
PRIOR FILING DATE: 1999-10-19
PRIOR APPLICATION NUMBER: 60/104,949
PRIOR APPLICATION NUMBER: 60/104,949
PRIOR FILING DATE: 1998-10-20
                                                                                                                                                       SOFTWARE: PatentIn Ver.
                                                                                                                                                                               NUMBER OF SEQ ID NOS: 13
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OTHER INFORMATION: May be any amino acid.
NAME/KEY: UNSURE
LOCATION: (234)
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OTHER INFORMATION: Description of Artificial Sequence: Kabat
OTHER INFORMATION: Consensus
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                                                                 ORGANISM: Artificial Sequence
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                                              EATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            59 YYSDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGT------YFFDSW 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYIS--SGGSGT 58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSVISGKTDGGST
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73.8%;
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Pred. No. 3.7e-40;
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                                                                                                    Query Match
Best Local S
Matches 96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
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APPLICANT: Whitlow, Marc D.
TITLE OP INVENTION: NO. 6764853el Method for Targeted Delivery of Nucleic Acids
FILE REFERENCE: 0977.2300002
CURRENT APPLICATION NUMBER: US/09/983,580
CURRENT FILING DATE: 2001-10-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: (234)
OTHER INFORMATION: May be a
NAME/KEY: UNSURE
LOCATION: (239)
OTHER INFORMATION: May be a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 13
SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1999-10-19
PRIOR APPLICATION NUMBER: 60/104,949
                                                                                                                                                                                                                                            LOCATION: (232)
LOCATION: (232)
OTHER INFORMATION: May be any amino NAME/KEY: UNSURE
LOCATION: (234)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1998-10-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 09/420,592
                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Descripti
OTHER INFORMATION: Consensus
NAME/KEY: UNSURE
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OTHER INFORMATION: May be NAME/KEY: UNSURE
                                                                                                                                                                                                                                           LOCATION: (234)
OTHER INFORMATION: May
                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                         OTHER INFORMATION: May be any amino acid
                                                                                                                                                                                                                       NAME/KEY: UNSURE
                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                            LOCATION: (239)
                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
                                  130 EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSVISGKTDGGST 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              110 GOGTLLTVSS
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59 YYSDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGT------YFDSW 109
                                                                                                                                                                                                                                                                                                                                                                                                                                          283
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                                                                   1 EVOLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYIS--SGGSGT
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| 96; Conservative
                                                                                                                    Similarity
                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , David R.
                                                                                                                                                                                                                                                                                                                                                                   Description of Artificial Sequence: Kabat
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73.8%;
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                                                                                                                    Score 485.5; DB 4
Pred. No. 3.7e-40;
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Pred. No. 3.7e-40;
9; Mismatches 14
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GENERAL INFORMATION:
APPLICANT: LAROSA, Gregory J.
APPLICANT: LAROSA, Christopher
APPLICANT: Newman, Christopher
APPLICANT: Newman, Walter
APPLICANT: O'Refe, Sidaran
APPLICANT: O'Refe, Theresa
TITLE OF INVENTION: HUMANIZED ANTI-CCR2 ANTIBODIES AND
TITLE OF INVENTION: METHODS OF USE THEREFOR
FILE REFERENCE: 1855.1052-004
CURRENT APPLICATION NUMBER: US/09/497,625A
CURRENT FILING DATE: 2000-02-03
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US-09-840-459-82
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US-09-497-625A-82
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CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: PCT/US01/03537
PRIOR FILING DATE: 2001-02-02
PRIOR PELICATION NUMBER: 09/497,625
PRIOR APPLICATION NUMBER: 09/497,625
PRIOR APPLICATION NUMBER: 09/359,193
PRIOR APPLICATION NUMBER: 09/359,193
PRIOR FILING DATE: 1999-07-22
PRIOR APPLICATION NUMBER: 09/121,781
PRIOR APPLICATION NUMBER: 09/121,781
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 107
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                              Sequence 82, Application US/09497625A Patent No. 6727349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local :
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LENGTH: 123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Horvath, Christopher
APPLICANT: Newman, Walter
APPLICANT: Jones, S. Tarran
APPLICANT: O'Brien, Siobhan H.
APPLICANT: O'Keefe, Tharesa
APPLICANT: O'Keefe, Tharesa
TITLE OF INVENTION: HUMANIZED ANTI-CCR2 ANTIBODIES AND
TITLE OF INVENTION: METHODS OF USE THEREFOR
FILE REFERENCE: 1855,1052-012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: LaRosa, Gregory J. APPLICANT: Horvath, Christop
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKKPGDYGSGSYYLDYWGQGTLVT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EVQLLESGGGLVQFGGSLRLSCAASGFTFSSYAMSWVRQAFGKGLEWVSAISGSGYTTYY
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76.4%;
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Pred. No. 1.6e-40;
9; Mismatches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16; Indels
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RESULT 33
US-08-983-607-32
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SOFTWARE: FRStSEQ for Windows Version
SEQ ID NO 82
LENGTH: 123
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Patent No. 61%.
Alan C
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                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/IB96
APPLICATION UNMBER: PCT/IB96
PILING DATE: June 28, 1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Mary M. Krinsky
REGISTRATION NUMBER: 32423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 09/121,781 PRIOR FILING DATE: 1998-07-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 09/359,193 PRIOR FILING DATE: 1999-07-22
                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 140 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 06520-8114

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" 1.44 Mb diskette
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC
OPERATING SYSTEM: MS DOS
SOFTWARE: Word Processing
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                                    REFERENCE/DOCKET NUMBER: OC
TELECOMMUNICATION INFORMATION:
TELEPHONE: 203-773-9544
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/983,607 FILING DATE: April 27, 1998 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 266 Whi
CITY: New Haven
STATE: Connecti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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STRANDEDNESS:
                    TYPE:
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            l: 140 residues amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Connecticut
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      266 Whitney Avenue
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                                                                                       203-773-1183
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alan Garen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Department of Molecular Biophysics and Biochemistry, Yale University
single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human Anti-Tumor Monoclonal Anti-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             77.7%;
76.4%;
                                                                                                                                                                                                                                                    PCT/IB96/01032
                                                                                                                                             OCR-679
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Pred. No. 1.6e-40;
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RESULT 34
US-08-974-899-6
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US-08-974-899-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                   CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 60/031971
FILING DATE: 11/27/96
ATTORNEY AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHNONE: 650/225-1994
                                                                                                TELEFAX: 650/952-9881 INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/974,899
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens (melanoma patient immu-
ORGANISM: nized with autologous tumor cells)
INDIVIDUAL ISCHATE: peripheral blood lymphocytes
IMMEDIATE SOURCE:
LIBRARY: DM414 scFv antibodies obtained from
LIBRARY: fUSE5 fusion phage construct
CLONE: V86
                                  SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Jardieu, Paula M.
TITLE OF INVENTION: Humanized Anti-CD11a Antibodies
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Presta, Leonard G. APPLICANT: Jardieu, Paula M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORIGINAL SOURCE
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DESCRIPTION:
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                                                                                                                                           Matches
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GENERAL INFORMATION:
                                                                                                                                                           Query Match
Best Local
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Best Local (
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                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Winbatin (Genencech)
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/795,798
FILING DATE: 28-Feb-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: P10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
                                                                                                                                                                                                                               TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Genente
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jardieu, Paula M.
TITLE OF INVENTION: Humanized Anti-CD11a Antibodies
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61
                    61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119
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                                                                                        1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                     EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSVISGDGGSTYY
                                                                                                                                                                                                                                                                   LENGTH: 113 amino acids
TYPE: Amino Acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: California
                                                                                                                                                                                                                                                                                                                                          TELEFAX: 650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94080
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                                                                                                                                           Conservative
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79.0%;
                                                                                                                                                          77.1%;
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                                                                                                                                         Score 481; DB 4; Length 113; Pred. No. 3.7e-40; 6; Mismatches 13; Indels
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Pred. No. 3.7e-40;
6; Mismatches 13
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                                                                                                                                           Gaps
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113
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RESULT 36 US-09-025-769B-38

Sequence 38, Application US/09025769B Patent No. 6300064

GENERAL INFORMATION:

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US-09-025-769B-63
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                                                                                                                                                                                                                        Sequence 63, Application US/09025769B Patent No. 6300064
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/025,769B
FILING DATE: 18-FEB-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 95 11 3021.0
FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: James F. Haley, Jr., Esq.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: MORPHO/5
                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Knappi
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                                                                                                                                              APPLICANT:
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TOPOLOGY: linear
MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
CORRESPONDENCE ADDRESS:
                                TITLE OF INVENTION: Protein/(Poly)peptide libraries NUMBER OF SEQUENCES: 373
CORRESPONDENCE ADDRESS:
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                                                                                                                               APPLICANT:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
ADDRESSEE: James F. Haley, Jr., Esq. c/o Fish & Neave STREET: 1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95;
                                                                                                                                                                                                                                                                                                                                      61 ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARWGGDGFYAMDYWGQGTLVTVSS
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    James F. Haley, Jr., Esq. c/o Fish & Neave
1251 Avenue of the Americas

                                                                                                                                             Knappik, Achim
Pack, Peter
Ilag, Vic
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Moroney, Simon
Plueckthun, Andreas
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oney, Simon
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79.2%;
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Pred. No. 4.4e-40;
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Patent No. 6696248
GENERAL INFORMATION:
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (212)596-9090
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: EP 95 11 3021.0
APPLICATION NUMBER: EP 95 11 3021.0
FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: James F. Haley, Jr., Esq.
REGISTRATION NUMBER: 27,794
REGISTRATION NUMBER: 27,794
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 10021
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/0: FILING DATE: 18-FEB-1998
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                                                     COMPUTER RADABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/490,070A
                                                                                                                                                                                                                                                                                                                                               Moroney, Simon
Plueckthun, Andreas
PITILE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
FILING DATE: 24-Jan-2000 PRIOR APPLICATION DATA: APPLICATION NUMBER: EP 9
                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
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95; Conserv
                                                                                                                                                                                          STATE: D.C.
COUNTRY: USA
ZIP: 20006
                                                                                                                                                                                                                                                  STREET: 1666 K Street, N.W., Suite 300 CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARWGGDGFYAMDYWGQGTLVTVSS
                                                                                                                                                                                                                                                                                        ADDRESSEE: Colin G. Sandercock, Esq. c/o Heller Ehrman White & McAuliffe
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79.2%;
   NUMBER: EP 95 11 3021.0
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Pred. No. 4.4e-40;
                                                                                               Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 3;
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                                                                                                 #1.30
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FILING DATE: 18-AUG-1995

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US-09-490-070A-63
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GENERAL INFORMATION:
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                       INFORMATION FOR SEQ ID NO: 63: SEQUENCE CHARACTERISTICS:
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                                                                                 APPLICATION NUMBER: EP 95 11 3021.0
PILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: Colin G. Sandercock, Esq.
REGISTRATION NUMBER: 31,298
REGISTRATION NUMBER: 37629-0005
TELEPONUNICATION INFORMATION:
TELEPHONE: (202) 912-2000
                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20006
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/490,070A FILING DATE: 24-Jan-2000 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
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TOPOLOGY: linear
MOLECULE TYPE: protein
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NAME: COlin G. Sandercock, Esq.
REGISTRATION NUMBER: 31,298
REFERENCE/DOCKET NUMBER: 37629-0005
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 912-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Colin G. Sandercock, Esq. c/o Heller Ehrman White & McAullffe
STREET: 1666 K Street, N.W., Suite 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EVOLVESGGGLVQPGGSLRLSCAASGPTPSSYAMSWVRQAPGKGLEWVSAISGSGGSTYY
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Ilag, Vic
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120 amino acids
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TOPOLOGY: linear;
; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 38:
US-09-490-153-38
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US-09-490-153-38
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Query Match
Best Local Similarity 79.2
                                                                                                                                                                                                         APPLICATION NUMBER: US/09/025,769B
FILING DATE: 18-FEB-1998
APPLICATION NUMBER: EP 95 11 3021.0
FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
NAME: James F: Haley, Jr., Esq.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: MORPHO/5
TELECHOME (212)596-9000
TELEPHONE: (212)596-9090
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESSEE: James F. Haley, Jr., Esq. c/o Fish & Neave ATREET: 1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 24-Jan-2000 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/490,153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Protein/(Poly)peptide libraries
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                                                                                                                                                      LENGTH: 120 amino acids
TYPE: amino acid
STRANDEDNESS: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 10021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: New York
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Plueckthun, Andreas
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Ilag, Vic
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             77.0%; Score 480.5; DB 4
79.2%; Pred. No. 4.4e-40;
tive 6; Mismatches 18
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79.2%;
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Pred: No. 4.4e-40;
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               Indels
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Search Job tim	рb	γQ	Ф	Ş
Search completed: December 17, 2004, 18:32:11 Job time : 73.2022 secs	61 ADSVKGRFTISRDNSKNTLYLÓMNSLRÁEDTAVYYCARWGGDGFYAMDYWGQGTLVTVSS 120	61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTR-VKLGTYYFDSWGQGTLLTVSS 119	1 EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYY 60	1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYI8SGGSGTYY 60

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Result
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Listing
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1: /cgn2_6/ptodata/2/pubpaa/US07_2: /cgn2_6/ptodata/2/pubpaa/PCT_N
3: /cgn2_6/ptodata/2/pubpaa/US06_
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                                                                                                                                                                                                                                                                                                                                                                                                                  / Cgm2 6/ptcodata/2/pubpaa/US09B_UBCOMB.pep:*
/ Cgm2 6/ptcodata/2/pubpaa/US09C_PUBCOMB.pep:*
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/ Cgm2 6/ptcodata/2/pubpaa/US10_NEW_PUB.pep:*
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                      US-09-811-123-3
4 US-10-268-501-6
5 US-10-608-626-6
5 US-10-600-152-15
6 US-10-619-754-6
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4 US-10-265-713-8
4 US-10-166-626-8
4 US-10-044-896-6
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US-10-166-626-18
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Sequence 18, Appl
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Sequence 15, Appli
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Sequence 6, Appli
Sequence 4, Appli
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0-474-	-10-693-62	-10-040-244-	-10-737-	-10-251-08	-10-474-832-6	2-6	-10-800-197-	-10-293-418-1	-09-880-748-14	-10-733-563-8	0-766-610-	-10-766-773-8	US-09-840-459-82	-09-985-442-	-09-983-580	-09-956-087-	-09-956-086-	US-10-835-641-21	-09-837-306	-10-045-674	US-10-045-674-487	-09-837-306	US-10-293-418-1953	80-748-195	US-10-269-805-59	-7	US-09-917-410-4	-10-128-520-	0-469-304-1	-10-291-265-	US-10-128-520-148
4, 1	e 64,	16,	e 93,	e 93,	e 69,	e 68,	e 1	142	_	8	8	е В	e 82	e 6,	9	w	w	Sequence 21, Appl			e 48	e 184,	195	195	Sequence 59, Appl	78,	, Appli	equence 156, Ap	17, App	equence 427,	Sequence 148, App

## ALIGNMENTS

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CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER FILING DATE: 1999-01-05
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER FILING DATE: 1995-03-21
EARLIER FILING DATE: 1995-03-21
EARLIER FILING DATE: 1996-03-21
EARLIER FILING DATE: 1997-09-17
EARLIER FILING DATE: 1994-09-17
EARLIER APPLICATION NUMBER: US07/947,674
EARLIER APPLICATION NUMBER: US08/408,1333
EARLIER 
; FEATURE:
; OTHER INFORMATION: cDNA KM-641
US-09-764-304-18
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US-09-764-304-18
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APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
APPLICANT: KUWANA, YOSHIHISA
APPLICANT: KUWANA, YOSHIHISA
APPLICANT: KUWANA, YOSHIHISA
FILE REFERENCE: 249-101
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                                                                                                                                                                      LENGTH: 130
TYPE: PRT
ORGANISM: Artificial Sequence
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APPLICANT: HASEGAWA, MAMORU
APPLICANT: MIXAAI, HIROMASA
APPLICANT: KUMANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HU
FILE REFERENCE: 249-101
CURRENT APPLICATION NUMBER: US/10/265,713
CURRENT FILING DATE: 2002-10-08
PRIOR APPLICATION NUMBER: US/09/225,322
PRIOR FILING DATE: 1995-01-05
PRIOR FILING DATE: 1995-05-31
PRIOR FILING DATE: 1995-05-31
PRIOR PRILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-03-21
PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR APPLICATION NUMBER: US 07/947,674
PRIOR APPLICATION NUMBER: US07/947,674
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                                                                                                                                                                                                                                                                       US-10-166-626-18
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                                                                                                                                                                              Sequence 18, Application US/10166626
Publication No. US20030166876A1
GENERAL INFORMATION:
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Best Local
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Publication No.
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Best Local
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APPLICANT: SHITARA, KENYA
APPLICANT: HANAI, NOBUO
APPLICANT: HASEGAWA, MAWORU
APPLICANT: MIYAJI, HIROMASA
APPLICANT: MIYAJI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SHITARA, KENYA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Artificial Sequence
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Pred. No. 1e-44;
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Pred. No. 1e-44;
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; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: CDNA KM-641
US-09-764-304-8
                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/764,304
CURRENT FILING DATE: 2001-01-19
EARLIER APPLICATION NUMBER: 09/225,322
EARLIER FILING DATE: 1999-01-05
EARLIER APPLICATION NUMBER: US 08/454,680
EARLIER FILING DATE: 1995-05-31
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER APPLICATION NUMBER: US 08/408,133
EARLIER FILING DATE: 1995-03-21
EARLIER APPLICATION NUMBER: US 08/292,178
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                                                                                                                                   SOFTWARE: PatentIn
SEQ ID NO 8
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Best Local Similarity
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PRIOR FILING DATE: 1991-09-18
NUMBER OF SEQ ID NOS: 19
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APPLICANT: HANAI, NOBUO
APPLICANT: HASECAWA, MAMORU
APPLICANT: MIYANI, HIROMASA
APPLICANT: KUWANA, YOSHIHISA
TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
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PRIOR FILING DATE: 1999-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
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CURRENT APPLICATION NUMBER: US/10/166,626
CURRENT FILING DATE: 2002-06-12
                                                                                                                                                                                                   EARLIER APPLICATION NUMBER: US07/947,674
EARLIER FILING DATE: 1992-09-17
EARLIER APPLICATION NUMBER: JP 3-238375
EARLIER FILING DATE: 1991-09-18
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PRIOR FILING DATE: 1995-03-21
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1994-08-17
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Pred. No. le-44;
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; Sequence 8, Application US/1016626; Publication No. US20030166876A1; GENERAL INFORMATION:
; APPLICANT: SHITARA, KENYA; APPLICANT: HANAI, NOBUO; APPLICANT: HANAI, NOBUO; APPLICANT: MIXEGAWA, MAMORU; APPLICANT: MIXEGAWA, MAMORU; APPLICANT: KUWANA, YOSHIHISA
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CURRENT FILING DATE: 2002-10-08
PRIOR APPLICATION NUMBER: US/09/225,322
PRIOR FILING DATE: 1999-01-05
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/454,680
PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR FILING DATE: 1995-03-21
PRIOR FILING DATE: 1994-08-17
PRIOR APPLICATION NUMBER: US 08/292,178
PRIOR FILING DATE: 1994-08-17
PRIOR APPLICATION NUMBER: US 07/947,674
PRIOR APPLICATION NUMBER: US 3-238375
PRIOR FILING DATE: 1992-09-18
PRIOR FILING DATE: 1991-09-18
PRIOR FILING DATE: 1991-09-18
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Best Local Similarity
Matches 108; Conser
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SOFTWARE: Patentin Ver. 2
SEQ ID NO 8
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TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY
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ORGANISM: Artificial Sequence
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MIYAJI, HIROMASA
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90.8%; Pred. No. 8
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Pred. No. 8.2e-44;
5; Mismatches 6
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-044-896-6
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NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 108; Conserv
                                                                                                                                                                    Matches
                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Kim, Jin K.
APPLICANT: Stewart, Timothy
APPLICANT: Presta, Leonard G.
TITLE OF INVENTION: ANTI-INTERPERON-ALPHA ANTIBODIES
FILE REFERENCE: GENENT.074A
CURRENT APPLICATION NUMBER: US/10/044,896
CURRENT FILING DATE: 2002-01-09
PRIOR APPLICATION NUMBER: 60/270775
PRIOR FILING DATE: 2001-02-22
NUMBER OF SEQ ID NOS: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Chuntharapai, Anan APPLICANT: Kim, Jin K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/408,133
PRIOR FILING DATE: 1995-03-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US/09/225,322B
PRIOR FILING DATE: 1999-01-05
PRIOR APPLICATION NUMBER: US 08/454,680
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/166,626
CURRENT FILING DATE: 2002-06-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: PROCESS FOR PRODUCING HUMANIZED CHIMERA ANTIBODY FILE REFERENCE: 249-101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1994-08-17
PRIOR APPLICATION NUMBER: USO7/947,674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 08/292,178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 130
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 1992-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     108; Conservative
61
                        61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119
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                                                                                                                                                                                                                                                                                                                                                          FastSEQ for Windows Version
                                                                                                         EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY 60
                                                                                 EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVAVISGDGGSTYY 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SDSVKGRFTISRDNAKNTLYLOMRSLRSEDSAMYFCTRVKLGTYYFDSWGQGTTLTVSS
  ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARGRVG--YYDYWGQGTLVTVSS
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90.8%;
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Pred. No. 1.6e-37;
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US-09-811-123-3

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CURRENT APPLICATION NUMBER: US/09/811,123
CURRENT FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/238,327
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 09/602,530
PRIOR FILING DATE: 2000-06-23
PRIOR FILING DATE: 2000-06-23
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FRAETSEQ for Windows Version 4.0
SEQ ID NO 3
SEQ ID NO 3
LENGTH: 119
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                                                                                                                                                                     US-10-268-501-6
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Publication No. US20030086924A1
GENERAL INFORMATION:
APPLICANT: Sliwkowski, Mark X.
TITLE OF INVENTION: Treatment with Anti-ErbB2 Antibodies
FILE REFERENCE: P1467R2P1
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Patent No. US20020001587A1
GENERAL INFORMATION:
APPLICANT: Sharon Erickson
APPLICANT: Ralph Schwall
APPLICANT: Mark Sliwkowski
                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 13
SEQ ID NO 6
LENGTH: 119
                                                                                 Matches
                                                                                                                            Query Match
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Best Local :
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CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US 09/602,812
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/141,316
PRIOR FILING DATE: 1999-06-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: METHODS OF TREATMENT USING ANTI-EXB
TITLE OF INVENTION: ANTIBODY-MAYTANSINOID CONJUGATES
FILE REFERENCE: GENENT.073A2
                                                                                                                                                                                    OTHER INFORMATION: heavy chain consensus sequence
                                                                                                                                                                                                            ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                                                                                                     TYPE: PRT
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                                                                                                         Local
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                       1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY 60
                                                                                                         Similarity
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EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVAVISGDGGSTYY
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                                                                                                79.8%;
79.8%;
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79.8%;
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                                                                               Score 498; DB 14;
Pred. No. 1.9e-37;
8; Mismatches 16
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Pred. No. 1.
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.9e-37;
                                                                                                                        DB 14;
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                                                                                 Indels
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CURRENT FILING DATE: 2003-06-27
PRIOR APPLICATION NUMBER: US 10/268,501
PRIOR FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US 09/602,812
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/141,316
PRIOR APPLICATION NUMBER: US 60/141,316
PRIOR FILING DATE: 1999-06-25
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                                                                                      US-10-600-152-15
                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/600,152
CURRENT FILING DATE: 2003-06-20
PRIOR APPLICATION NUMBER: 09/648,067
PRIOR FILING DATE: 2000-08-25
PRIOR FILING DATE: 1099-08-27
PRIOR FILING DATE: 1999-08-27
PRIOR FILING DATE: 1999-08-27
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: US 60/213,822
PRIOR FILING DATE: 2000-06-23
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Publication No. US20040037824A1
GENERAL INFORMATION:
APPLICANT: Baughman, Sharon A.
APPLICANT: Shak Steven
Query Match
Best Local Sim
Matches 95;
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SEQ ID NO 15
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Best Local
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SEQ ID NO 6
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APPLICANT: Sliwkowski, Mark X.
TITLE OF INVENTION: Treatment with Anti-ErbB2 Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Dosages for Treatment with Anti-ErbB2 Antibodies FILE REFERENCE: P1775R1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: P1467R2P2
                                                                                                                                        TYPE: PRT ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                FEATURE:
                                                                                                          OTHER INFORMATION: VH consensus sequence
                                                                                                                                                                                             LENGTH: 119
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                      Similarity
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    Conservative
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  79.8%; Score 498; DB 15; Length 119; 79.8%; Pred. No. 1.9e-37; tive 8; Mismatches 16; Indels
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Pred. No. 1.9e-37;
8; Mismatches 16;
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US-10-835-641-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Kelsey, Stephen
TITLE OF INVENTION: Methods for Identifying Tumors That Are
TITLE OF INVENTION: Responsive To Treatment With Anti-ErbB2 Antibodies
FILE REFERENCE: 39766-0114A
CURRENT APPLICATION NUMBER: US/10/619,754
CURRENT FILING DATE: 2003-07-14
PRIOR APPLICATION NUMBER: US 60/396,290
PRIOR FILING DATE: 2002-07-15
PRIOR FILING DATE: 2002-07-15
PRIOR APPLICATION NUMBER: US 60/480,043
PRIOR FILING DATE: 2003-06-20
NUMBER OF SEQ ID NOS: 6
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Koll, Hans
APPLICANT: Bossenmaier, B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 6, Application US/10619754
Publication No. US20040106161A1
                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/10835641 Publication No. US20040236078A1
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Best Local Similarity
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ORGANISM: Homosapiens
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                                                                                                                                                                                                                                                                         APPLICANT: Carter, Paul J.

Presta, Leonard G.
TITLE OF INVENTION: Method for Making Humanized Antibodies
            ZIP: 94080

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: WinPatin (Genentech)
                                                                                                                                                                                            NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 1 DNA Way
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95;
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                                                                                                                                                        CITY: South San Francisco
STATE: California
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                                                                                                                                     COUNTRY: USA
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Muller, Hans-Joachim
Sliwkowski, Mark
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Pred. No. 1.9e-37;
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US-10-128-520-148
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                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 381
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 148
LENGTH: 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Publication No. US20040105862A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local S
                                                                                                                                                                                                                                                                               Query Match
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CURRENT FILING DATE: 2002-04-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: PAN et al.
TITLE OF INVENTION: Human TIMP-1 Antibodies
FILE REFERENCE: 02973.00073
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/285,683 PRIOR FILING DATE: 2001-04-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
                                                                                                                                                                                                                                                             Local
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APPLICATION NUMBER: US/09/705,398

FILING DATE: 02-NOV-2000

APPLICATION NUMBER: 08/146206
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: P0709P1D2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 17-NOV-1993
APPLICATION NUMBER: 07/715272
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
119 88 120
                                     118 SS 119
                                                                                                                                                                                                                                      al Similarity
97; Conserva
                                                                           61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAR--LDTYYPDLFDYWGQGTLVTV 118
                                                                                                      13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTY-YFDSWGQGTLLTVSS
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                                                                                                                                                          1 QVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYY
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Similarity 80.8%;
97; Conservative
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TYPE: Amino Acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/10/835,641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/10128520
                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                          79.1%;
79.5%;
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Pred. No. 3.3e-37;
7; Mismatches 15
                                                                                                                                                                                                                                        Score 493.5; DB 16;
Pred. No. 9.4e-37;
7; Mismatches 13;
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US-10-291-265-427

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APPLICANT: Hyseq, Inc.
APPLICANT: Tang et al
ITITLE OF INVENTION: No. US20030232054Alel Nucleic Acids and Pol)
FILE REFERENCE: 21272-017 (785)
CURRENT APPLICATION NUMBER: US/10/291,265
CURRENT FILING DATE: 2000-01-25
PRIOR APPLICATION NUMBER: 09/491,404
PRIOR APPLICATION NUMBER: 09/491,404
PRIOR APPLICATION NUMBER: 09/617,746
PRIOR FILING DATE: 2000-01-25
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: 09/631,451
PRIOR APPLICATION NUMBER: 09/631,451
PRIOR APPLICATION NUMBER: 09/633,870
PRIOR APPLICATION NUMBER: 09/633,870
PRIOR FILING DATE: 2000-09-15
NUMBER OF SEQ ID NOS: 944
SOPTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 427
LENGTHH: 713
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US-10-291-265-427
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                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/469,304
CURRENT FILING DATE: 2003-08-29
PRIOR APPLICATION NUMBER: JP2001/317054
PRIOR FILING DATE: 2001-10-15
NUMBER OF SEQ ID NOS: 147
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 17
                                                                                                                                                                                                                                                                                                                                                                          Sequence 17, Application US/10469304
Publication No. US20040091974A1
GENERAL INFORMATION:
APPLICANT: KIRIN BEER KABUSHIKI KAISHA
TITLE OF INVENTION: Anti HLA-DR antibody
FILE REFERENCE: PH-1646-PCT
                                                                                                  Best
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Publication No.
                                                                              Matches
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Best Local :
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                                                                                                                   Query Match
                                                                                                                                                                                                   TYPE: PRT
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                                                                                                                                                                                                                    LENGTH: 143
                                                                                                Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     113 TLLTVSS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        121 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKSHPGYYYDSSGYSYYFDYWGQG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61
                    1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY 60
                                                                                96;
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                                                                                              Similarity
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EVQLLESGGGLVQPGGSLRLSCAASGFTFSSYAMTWVRQAPGKGLEWVSGISGGGDSTYY
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D. US20030232054A1
                                                                              Conservative
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74.8%;
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78.0%;
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                                                                            Score 489; DB 15;
Pred. No. 1.5e-36;
9; Mismatches 14;
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Pred. No. 1.8e-36;
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                                                                                                                 Length 143;
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CURRENT FILING DATE: 2002-04-24
PRIOR APPLICATION NUMBER: US 60/285,683
PRIOR FILING DATE: 2001-04-24
NUMBER OF SEQ ID NOS: 381
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/09917410 Patent No. US20020098183A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: PAN et al.
TITLE OF INVENTION: Human TIMP-1 Antibodies
FILE REFERENCE: 02973.00073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                  COMPUTER: IBM PC comparible operATING SYSTEM: PC-DOS/MS-DOS SOPTMARE: ASCII, WordPerfect 5.1 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/917,410 PILING DATE: 26-011-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Computer Disk
                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                       CO, Man S. TITLE OF INVENTION: A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: MARTIN, Ulrich; HASELBECK, Anton; SCHUMACHER, Guenther;
                  PRIOR
                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 140
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APPLICATION DATA: APPLICATION NUMBER: 08/578,953
                                                                                                                                                                                                                                      COUNTRY: U.S.A.
                                                                                                                                                                                                                                                          STATE: New York
                                                                                                                                                                                                                                                                         CITY: New York
                                                                                                                                                                                                                                                                                                 STREET: 805 Third Avenue
                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Felfe & Lynch
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                                                                                                                                                                                                                   10022
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  78.3%;
79.8%;
                                                                                                                                                                                                                                                                                                                                                                                                MULTIPLE ORGAN FAILURE AFTER POLYTRAUMA AND FOR PREVENTION OF ACUTE ORGAN DAMAGE AFTER
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                                                                                                                                                                                                                                                                                                                                                                             EXTRACORPOREAL BLOOD
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Pred. No. 2.6e-36;
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PRIOR APPLICATION NUMBER: 00110063.5
PRIOR FILING DATE: 2000-05-12
NUMBER OF SEQ ID NOS: 97
SOFTWARE: Patentin version 3.2
SEQ ID NO 78
LENGTH: 121
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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US-10-275-046-78
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APPLICANT: NAGY et al.
APPLICANT: NIVENTION: IMMUNOMODULATORY HUMAN MHC CLASS II ANTIGEN-BINDING POLYPEPTIDES
FILE REFERENCE: GPCG-P01-260
CURRENT APPLICATION NUMBER: US/10/275,046
CURRENT FILING DATE: 2002-10-31
CURRENT FILING DATE: 2002-10-31
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Best Local Similarity
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Best Local Similarity
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NAME: HARSON, NO. US20020099183Alman D.
REGISTRATION NUMBER: 30,946
REFERENCE/DOCKET NUMBER: BOER 1059-PFF/NDH/SLH
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
121 S 121
                                      119 S 119
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                                                                                                   61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTY--YFDSWGQGTLLTVS
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                                                                            61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARSSMPMYGEGFDLWGQGTLVTVS
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                                                                                                                                                     EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYY
                                                                                                                                                                                        EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PDSVKGRFTISRDNAKNTLYLOMNSLRAEDTAVYYCARDYDG--YFDYWGOGTLVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: double
TOPOLOGY: linear
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APPLICATION NUMBER: EP
FILING DATE: 19-Sep-95
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82.4%;
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Pred. No. 1.9e-36;
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Pred. No. 6.5e-36;
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RESULT 20

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; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 59
; LENGTH: 123
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-269-805-59
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                                                                                                                                                                                                                                         SOFTWARE: PatentIn Ver.
SEQ ID NO 1953
LENGTH: 247
TYPE: PRT
                                                                                                                                         Query Match
Best Local :
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Publication No. US20
GENERAL INFORMATION:
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Best Local Similarity
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TITLE OF INVENTION: ANGIOPOTETIN-2 SPECIFIC BINDING AGENTS
FILE REFERENCE: A-722
CURRENT APPLICATION NUMBER: US/10/269,805
CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US 60/328,604
PRIOR PLING DATE: 2001-10-11
NUMBER: OF SEQ ID NOS: 76
NUMBER OF SEQ ID NOS: 76
                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2001-05
NUMBER OF SEQ ID NOS: 3239
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CURRENT FILING DATE: 2001-06-15
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
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                                                                                                                                                                                                                      ORGANISM: Homo sapiens
                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 60/276,248
FILING DATE: 2001-03-16
APPLICATION NUMBER: 60/277,379
FILING DATE: 2001-03-21
                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/293,499
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94;
                                                                              1 EVOLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY 60
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                                          EVOLVESGGGLVQPGWSLRLSCAASGFTFSYSAMTWVRQAPGKGLEWVSAITSSGGATYY 60
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No. US20030124129A1
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                                                                                                                      Conservative
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                                                                                                                                         78.0%; Score 487; DB 1 76.5%; Pred. No. 4e-36;
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                                                                                                                                                            DB 10; Length 247;
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US-09-837-306-184
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US-10-293-418-1953
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PRIOR APPLICATION NUMBER: 60/331,469
PRIOR PELING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR PILING DATE: 2001-12-19
PRIOR PILING DATE: 2001-06-15
PRIOR PILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR PILING DATE: 2001-03-16
PRIOR PILING DATE: 2000-10-17
PRIOR PILING DATE: 2000-10-17
                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: LADNER, ROBERT C.
APPLICANT: COHEN, EDWARD H.
APPLICANT: NASTRI, HORACIO G.
APPLICANT: ROOKEY, KRISTIN L.
                APPLICANT: ROCKEY, KRISTIN L.

APPLICANT: HOET, RENE
TITLE OF INVENTION: NOVEL METHODS OF CONSTRUCTING LIBRARIES OF GENETIC
TITLE OF INVENTION: PACKAGES THAT COLLECTIVELY DISPLAY THE MEMBERS OF A
TITLE OF INVENTION: DIVERSE FAMILY OF PEPTIDES, POLYPEPTIDES OR PROTEINS
FILE REFERENCE: DYAX/002
CURRENT APPLICATION NUMBER: US/09/837,306
CURRENT PILING DATE: 2001-09-24
PRIOR APPLICATION NUMBER: 60/198,069
PRIOR PEPLICATION NUMBER: 60/198,069
PRIOR FILING DATE: 2000-04-17
NUMBER OF SEQ ID NOS: 428
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 184
LENGTH: 136
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LENGTH: 247
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Publication No. US20
GENERAL INFORMATION
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Best Local (
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CURRENT APPLICATION NUMBER: US/10/293,418
CURRENT FILING DATE: 2002-11-27
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TYPE: PRT
TYPE:
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APPLICATION NUMBER: 60/212,210
FILING DATE: 2000-06-16
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Similarity 76.5%;
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RESULT 25
US-10-045-674-453
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Sequence 453, Application US/10045674
Publication No. US20030232333A1
GENERAL INFORMATION:
APPLICANT: LADNER, ROBERT C.
APPLICANT: COHEN, EDWARD H.
APPLICANT: NASTRI, HORACIO G.
APPLICANT: ROOKEY, KRISTIN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 09/837,306
PRIOR FILING DATE: 2001-04-17
NUMBER OF SEQ ID NOS: 635
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: LADNER, ROBERT C.
APPLICANT: COHEN, EDWARD H.
APPLICANT: NASTRI, HORACIO G.
APPLICANT: ROOKEY, KRISTIN L.
                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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Best Local
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TITLE OF INVENTION: NOVEL METHODS OF CONSTRUCTING LIBRARIES COMPRISING
TITLE OF INVENTION: DISPLAYED AND/OR EXPRESSED MEMBERS OF A DIVERSE FAMI
TITLE OF INVENTION: OF PEPTIDES, POLYPEPTIDES OR PROTEINS AND THE NOVEL
TITLE OF INVENTION: LIBRARIES
FILE REFERENCE: DYAX/002 CIP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/045,674
CURRENT FILING DATE: 2001-10-25
PRIOR APPLICATION NUMBER: 60/198,069
PRIOR FILING DATE: 2000-04-17
PRIOR FILING DATE: 2000-04-17
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Unknown Organism
FEATURE:
OTHER INFORMATION: Description of Unknown Organism: V3-23
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                                                                                                                                                                                                                                                                                                                                                                      94;
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                                                                                                                                                                                                                     ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCAKDYEGTGYAFDIWGQGTMVTVSS 126
                                                                                                                                                                                                                                                                                               EVOLLESGGGLVOPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYY
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78.3%;
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Pred. No. 2.4e-36;
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Pred. No. 2.4
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CURRENT APPLICATION NUMBER: US/10/045,674
CURRENT FILING DATE: 2001-10-25
PRIOR APPLICATION NUMBER: 60/198,069
PRIOR FILING DATE: 2000-04-17
PRIOR APPLICATION NUMBER: 09/837,306
PRIOR FILING DATE: 2001-04-17
NUMBER OF SEQ ID NOS: 635
SOFTWARE: PACENTIN Ver. 2.1
SEQ ID NO 453
LENGTH: 367
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                                                                                                                                                                             US-09-837-306-196
                                                                                                                                                                                                                                                                             TITLE OF INVENTION: NOVEL METHODS OF CONSTRUCTING LIBRARIES OF GENETIC TITLE OF INVENTION: PACKAGES THAT COLLECTIVELY DISPLAY THE MEMBERS OF A TITLE OF INVENTION: DIVERSE FAMILY OF PEPTIDES, POLYPEPTIDES OR PROTEINS PLE REFERENCE: DYAX/002

CURRENT PELLORION NUMBER: US/09/837,306

CURRENT FILING DATE: 2001-09-24

PRIOR APPLICATION NUMBER: 60/198,069

PRIOR APPLICATION UMBER: 60/198,069

PRIOR FILING DATE: 2000-04-17

NUMBER: OF SEQ ID NOS: 428

SOFTWARE: PACENTIN Ver. 2.1

SEQ ID NO 196

LENGTH: 368
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                                                                                         Matches
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APPLICANT: CCHEN, EDWARD H.
APPLICANT: NASTRI, HORACIO G.
APPLICANT: ROOKEY, KRISTIN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: HOST, RENE
APPLICANT: HOOGENBOOM, HENDRICUS R. J. M.
TITLE OF INVENTION: NOVEL METHODS OF CONSTRUCTING LIBRARIES COMPRISING
TITLE OF INVENTION: DISPLAYED AND/OR EXPRESSED MEMBERS OF A DIVERSE FAMILY
TITLE OF INVENTION: OF PEPTIDES, POLYPEPTIDES OR PROTEINS AND THE NOVEL
TITLE OF INVENTION: LIBRARIES
FILE REFERENCE: DYAX/002 CIP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Unknown Organism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Unknown Organism: MALIA3 protein OTHER INFORMATION: sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Unknown Organism FEATURE:
                                                                                                                                                                                                FEATURE:
OTHER INFORMATION: Description
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nes 94; Conservative
                                                                                                             Local
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                         1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                           Similarity
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EVOLLESGGGLVOPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYY
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78.3%;
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                                                                                    Score 486.5; DB 1
Pred. No. 6.6e-36;
9; Mismatches 16
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Pred. No. 6.6e-36;
                                                                                                                                                                                           of Unknown Organism: MALIA3
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                                                                                                                             Length 368;
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                                                                                  Gaps
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RESULT 28
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Publication No. US2004
GENERAL INFORMATION:
                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 650/952-9881 INFORMATION FOR SEQ ID NO: 21:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/705,398
FILING DATE: 02-Nov-2000
APPLICATION NUMBER: 08/146206
FILING DATE: 17-NOV-1993
APPLICATION NUMBER: 07/715272
FILING DATE: 14-JUN-1991
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REGISTRATION NUMBER: 40,378
                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/835,641
FILING DATE: 30-Apr-2004
CLASSIFICATION: dTuknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                  TYPE: Amino acids
TYPE: Amino Acid
TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb |
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Presta, Leonard
TITLE OF INVENTION: Method
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION: 650/225-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Carter, Paul.J.
                                                  121
                                                                                   118 SS 119
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                                                                                                                                          83 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKDYEGTGYAFDIWGQGTMVTVSS 142
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                                                                                                                      ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGRVGYSLSGLYDYWGQGTLVTV 120
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COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 94080
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                                                    122
                                                                                                                                                                                                                                                                               77.8%;
77.0%;
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                                                                                                                                                                                                                                                           10; Mismatches
                                                                                                                                                                                                                                                                             Score 485.5; DB 1
Pred. No. 2.7e-36;
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for Making Humanized Antibodies
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US-09-956-086-3

Sequence 3, Application US/09956086 Patent No. US20020155498A1

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RESULT 29
US-09-956-087-3
; Sequence 3, Application US/09956087
; Patent No. US20020161201A1
; GENERAL INFORMATION:
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Best Local (
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APPLICANT: FILE
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APPLICATION NUMBER: 09/069,821
FILING DATE: <Unknown:
APPLICATION NUMBER: US 60/063,074
FILING DATE: 27-OCT-1997
APPLICATION NUMBER: US 60/050,472
FILING DATE: 23-JUN-1997
APPLICATION NUMBER: US 60/044,449
FILING DATE: 30-APR-1997
                                                                                                                                                                                                                                                                                                                                                                 Match 77.8%;
Local Similarity 73.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single
TOPOLOGY: No. US20020155498A1 Relevant
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide SEQUENCE DESCRIPTION: SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 263 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: KIM, JUDITH U.
REGISTRATION NUMBER: 40,679
REFERENCE/DOCKET NUMBER: 0977.2280003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: STERNE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
                                                                                                                                                                                                        190
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                                                                                                                                                                                                                                                                                               1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYIS--SGGSGT
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                                                                                                                                                                                                      YYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGRXGXSLSGXYYYYHYFDYW
                                                                                                                              GQGTLVTVSS
                                                                                                                                                                  GOGTLLTVSS 119
                                                                                                                                                                                                                                      YYSDSVKGRFTISRDNSKNTLYLOMRSLRAEDSAVYFCTRVKLGT-----YYFDSW 109
                                                                                                                                                                                                                                                                          EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSVISGKTDGGST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 1100 NEW YORK AVE., NW, SUITE 600 CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/956,086 FILING DATE: 20-Sep-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: DC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LEE, LIHSYNG S.
OF INVENTION: SINGLE-CHAIN ANTIGEN-BINDING PROTEINS
CAPABLE OF GLYCOSYLATION, PRODUCTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (202)371-2600
TELEFAX: (202)371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 20005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                259
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                                                                                                                                                                                                                                                                                                                                                                 Score 485.5; DB 9;
Pred. No. 5.8e-36;
                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                    Length 263;
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                                                                                                                                                                                                                                                                            189
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LENGTH: 263 amino acids
TYPE: amino acids
STRANDEDNESS: single
TOPOLOGY: not relevant
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 3:
US-09-956-087-3
RESULT 30
US-09-983-580-6
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Best Local Similarity
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TELEPAX: (202)371-2540
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 09/069,821
FILING DATE: 1998-04-30
APPLICATION NUMBER: US 60/063,074
FILING DATE: 27-0CT-1997
APPLICATION NUMBER: US 60/050,472
FILING DATE: 23-JUN-1997
APPLICATION NUMBER: US 60/044,449
FILING DATE: 30-APR-1997
APPLICATION NUMBER: US 60/044,449
FILING DATE: 30-APR-1997
ATTORNEY/AGENT INFORMATION:
RAME: KIM, JUDITH U.
REGISTRATION NUMBER: 40,679
REFERENCE/DOCKET NUMBER: 0977.228000
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin R CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.
STREET: 1100 NEW YORK AVE., NW, SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LEE, LIHSYNG S.
TITLE OF INVENTION: SINGLE-CHAIN ANTIGEN-BINDING PROTEINS
CAPABLE OF GLYCOSYLATION, PRODUCTION AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                  190
                                                                                                                                                                                                                                                 130 EVOLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSVISGKTDGGST 189
                                                                                    250
                                                                                                                        110 GOGTLLTVSS 119
                                                                                                                                                                                          59 YYSDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGT-------YYFDSW 109
                                                                                                                                                                                                                                                                        1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYIS--SGGSGT 58
                                                                                                                                                                                                                                                                                                                                 96;
                                                                                    GQGTLVTVSS
                                                                                                                                                                  YYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGRXGXSLSGXYYYYHYFDYW 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/956,087 FILING DATE: 20-Sep-2001 CLASSIFICATION: <Unknown>
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                                                                                                                                                                                                                                                                                                                                 Conservative
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SHORR, ROBERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
                                                                                                                                                                                                                                                                                                                                                  77.8%;
73.8%;
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Pred. No. 5.8e-36;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Version #1.30
                                                                                                                                                                                                                                                                                                                                                                     9;
                                                                                                                                                                                                                                                                                                                                                                     Length 263;
                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                   11;
                                                                                                                                                                                                                                                                                                                                 Gaps
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; Sequence 6, Application US/09983580 Patent No. US20020151061A1 ; GENERAL INFORMATION: APPLICANT: Filpula, David R.

Wang, Maoliang

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CURRENT APPLICATION NUMBER: US/09/983,580
CURRENT FILING DATE: 2001-10-25
PRIOR APPLICATION NUMBER: 09/420,592
PRIOR FILING DATE: 1999-10-19
PRIOR APPLICATION NUMBER: 60/104,949
PRIOR FILING DATE: 1998-10-20
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 6
LENGTH: 283
TYPE: PRT
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US-09-985-442-6
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                                                                                                                                                               GENERAL INFORMATION:

APPLICANT: Filpula, David R.
APPLICANT: Wang, Macliang
APPLICANT: Wang, Marc D.

TITLE OF INVENTION: No. US20020156248A1el Method for Targeted Delivery of Nucleic Aci
FILE REFERENCE: 0977.2300003
CURRENT FILING DATE: 2001-11-02
PRIOR APPLICATION NUMBER: US/09/985,442
CURRENT FILING DATE: 1999-10-19
PRIOR APPLICATION NUMBER: 09/420,592
PRIOR FILING DATE: 1999-10-19
PRIOR APPLICATION NUMBER: 60/104,949
PRIOR FILING DATE: 1988-10-20
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                               SEQ ID NO 6
LENGTH: 283
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TITLE OF INVENTION: No. US20020151061A1el Method for Targeted Delivery of Nucleic
FILE REFERENCE: 0977.2300002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: May be any amino acid.
NAME/KEY: UNSURE
LOCATION: (234)
OTHER INFORMATION: Description of Artificial Sequence: Kabat OTHER INFORMATION: Consensus 'NAME/KEY: UNSURE'
                                                                                     TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: May be
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                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         110 GOGTLLTVSS 119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYIS--SGGSGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   YYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGRXGXSLSGXYYYYHYFDYW 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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73.8%;
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Pred. No. 6.3e-36;
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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Horvath, Christopher

APPLICANT: Horvath, Christopher

APPLICANT: Jones, S. Tarran

APPLICANT: Jones, S. Tarran

APPLICANT: O'Brien, Siobhan H.

APPLICANT: O'Keefe, Theresa

TITLE OF INVENTION: HUMANIZED ANTI-CCR2 ANTIBODIES AND

TITLE OF INVENTION: METHODS OF USE THEREFOR

FILLE REFERENCE: 1855.1052-012

CURRENT APPLICATION NUMBER: US/09/840,459

CURRENT PILING DATE: 2001-02-02

PRIOR APPLICATION NUMBER: PCT/US01/03537

PRIOR APPLICATION NUMBER: 99/497,625

PRIOR APPLICATION NUMBER: 09/497,625


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                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 82
LENGTH: 123
                                                                                                                                               Matches
                                                                                                                                                                 Query Match
Best Local
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Best Local
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                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1998-07-23
                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 1999-07-22
PRIOR APPLICATION NUMBER: 09/121,781
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: May be any amino NAME/KEY: UNSURE LOCATION: (234)
OTHER INFORMATION: May be any amino NAME/KEY: UNSURE LOCATION: (239)
                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: May be
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US200,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  250 GOGTLVTVSS 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        190 YYADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARGRXGXSLSGXYYYYHYFDYW
5
                                 61 SDSVKGRFTISRDNSKNTLYLOMRSLRAEDSAVYFCTR----VKLGTYYFDSWGQGTLLT 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           59 YYSDSVKGRFTISRDNSKNTLYLOMRSLRAEDSAVYFCTRVKLGT------YYFDSW
                                                                                                                                               94;
                                                                                         1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                                                                                 Similarity
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ADSVKGRETISRDNSKNTLYLOMNSLRAEDTAVYYCAKKPGDYGSGSYYLDYWGQGTLVT
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S20020150576A1
                                                                                                                                               Conservative
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73.8%;
                                                                                                                                                               77.78;
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                                                                                                                                               9; Mismatches
                                                                                                                                                               Score 485; DB 9
Pred. No. 3e-36;
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Pred. No. 6.3e-36;
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                                                                                                                                                                              DB 9;
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                                                                                                                                                                                Length 123;
                                                                                                                                               Indels
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117

VSS 119

RESULT 33 US-10-766-773-82 뭐

VSS

123

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Sequence 82, Application US/10766610

Publication No. US20040132980A1

GENERAL INFORMATION:

APPLICANT: HORVATH, Christopher

APPLICANT: Horvath, Christopher

APPLICANT: Newman, Walter

APPLICANT: O'Brien, Siobhan H.

APPLICANT: O'Keefe, Theresa

TITLE OF INVENTION: HUMANIZED ANTI-CCR2 ANTIBODIES AND

TITLE OF INVENTION: HUMANIZED ANTI-CCR2 ANTIBODIES AND

TITLE OF INVENTION: WETHODS OF USE THEREFOR

FILE REFERENCE: 1855.1052-029

CURRENT APPLICATION NUMBER: US/10/766,610

CURRENT APPLICATION NUMBER: 09/840,459

PRIOR APPLICATION NUMBER: 001-02-02

PRIOR APPLICATION NUMBER: 0074-23

PRIOR APPLICATION NUMBER: 0074-23

PRIOR APPLICATION NUMBER: 009/840,459

PRIOR APPLICATION NUMBER: 009/840,459

PRIOR APPLICATION NUMBER: 009/840,459

PRIOR APPLICATION NUMBER: 0074-23

PRIOR APPLICATION NUMBER: 0074-23

PRIOR APPLICATION NUMBER: 0074-625

PRIOR APPLICATION NUMBER: 0074-625

PRIOR APPLICATION NUMBER: 0074-625
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FILE REFERENCE: 1855.1052-028
CURRENT APPLICATION UMBER: US/10/766,773
CURRENT FILING DATE: 2004-01-27
PRIOR APPLICATION NUMBER: 09/497,625
PRIOR APPLICATION NUMBER: 09/497,625
PRIOR APPLICATION NUMBER: 09/39,193
PRIOR APPLICATION NUMBER: 09/359,193
PRIOR FILING DATE: 1999-07-22
PRIOR FILING DATE: 1999-07-23
I PRIOR APPLICATION NUMBER: 09/121,781
PRIOR APPLICATION NUMBER: 09/121,781
PRIOR APPLICATION NUMBER: 09/121,781
PRIOR APPLICATION NUMBER: 09/121,781
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 106
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 82
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US-10-766-610-82
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TYPE: PRT
ORGANISM: Homo sapiens
US-10-766-773-82
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Publication No. US20040126851A1
GENERAL INFORMATION:
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APPLICANT: Horvath, Christophe:
APPLICANT: Newman, Walter
APPLICANT: Jones, S. Tarran
APPLICANT: O'Brien, Siobhan H.
APPLICANT: O'Keefe, Theresa
FILING DATE:
APPLICATION 1
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76.4%;
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Pred. No. 3e-3
9; Mismatches
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RESULT 36 US-09-880-748-1421

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CURRENT FILING DATE: 2003-12-10
PRIOR APPLICATION NUMBER: US 10/272,899
PRIOR FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: US 60/392,364
PRIOR FILING DATE: 2002-06-26
PRIOR APPLICATION NUMBER: US 60/350,166
PRIOR APPLICATION NUMBER: US 60/350,166
PRIOR PILING DATE: 2001-10-19
NUMBER OF SEQ ID NOS: 122
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; ORGANISM: Homo sapiens
US-10-766-610-82
                                                                                                                                                                                                                                                                            US-10-733-563-82
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Publication No. US20040151721A1
GENERAL INFORMATION:
APPLICANT: O'Keefe, Theresa
APPLICANT: Ponath, Paul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1999-07-22
PRIOR APPLICATION NUMBER: 09/121,781
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 107
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 82
                                                                                                                                                                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 82
                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 94; Conserv
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Best Local Similarity
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TITLE OF INVENTION: HUMANIZED ANTI-CCR2 ANTIBODIES AND
TITLE OF INVENTION: METHODS OF USE THEREOF
FILE REFERENCE: 10448-213001
                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                             TYPE: PRT
                                                                                                                                                                                                                                                                                                                                 LENGTH: 123
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                              117 VSS 119
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                                                                                    61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKKPGDYGSGSYYLDYWGQGTLVT
                                                                                                                                                         1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY 60
                                                                                                                                                                                                        94; Conservative
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VSS 123
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Pred. No. 3e-36;
                                                                                                                                                                                                         Score 485; DB 1
Pred. No. 3e-36;
9; Mismatches
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                                                                                                                                                                                                                                            DB 16; Length 123;
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RESULT 37
US-10-293-418-1421
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1421, Application US/10293418 Publication No. US20030223996A1 GENERAL INFORMATION:
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Best Local :
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             SEQ ID NO 1421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Ruben et al.

TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523
CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-15
PRIOR FILING DATE: 2000-06-15
                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/293,418
CURRENT FILING DATE: 2002-11-27
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILING DATE: 2001-12-19
PRIOR FILING DATE: 2001-12-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 3239
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PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/277,379
                                         NUMBER OF SEQ ID NOS:
                                                           PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-16
                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 09/880,748
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLySFILE REFERENCE: PF523P2
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ENGTH:
                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/293,499
                                                                                                    APPLICATION NUMBER: 60/240,816
FILING DATE: 2000-10-17
                                                                                                                                                                     APPLICATION NUMBER: 60/276,248
                                                                                                                                                                                                              APPLICATION NUMBER: 60/277,379
                                                                                                                                                                                                                                     FILING DATE: 2001-05-25
                                                                                                                                                  FILING DATE: 2001-03-16
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76.8%;
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Pred. No. 6.1e-36;
8; Mismatches 15;
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Sequence 68, Application US/10474832
Publication No. US20040081651A1
GENERAL IMPORMATION:
TITLE OF INVENTION: ANTIBODIES TO VLA-1
FILE REFERENCE: A101 PCT
CURRENT APPLICATION NUMBER: US/10/474,832
CURRENT EILING DATE: 2003-10-14
PRIOR APPLICATION NUMBER: 60/283,794
PRIOR APPLICATION NUMBER: 60/283,794
PRIOR APPLICATION NUMBER: 60/303,689
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Best Local S
Matches 93
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15, Applica Publication No. US20 GENERAL INFORMATION:
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Best Local Similarity 76.8%;
Matches 96; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/800,197
CURRENT FILING DATE: 2004-03-12
PRIOR APPLICATION NUMBER: 60/455,094
PRIOR FILING DATE: 2003-03-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Morton, Philip A et
TITLE OF INVENTION: ANTIBODIES
FILE REFERENCE: 01343/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: phage display generated antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
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No. US20040202655A1
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Pred. No. 6.1e-36;
8; Mismatches 15
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WS-10-474-832-69
US-10-474-832-69

US-10-474-832-69
; Sequence 69, Application US/10474832
; Publication No. US20040081651A1
; GENERAL INFORMATION:
; APPLICANT: BICGEN, INC.
; TITLE OF INVENTION: ANTIBODIES TO VLA-1
; FILE REFERENCE: A101 PCT
; CURRENT APPLICATION NUMBER: US/10/474,832
; CURRENT FILING DATE: 2003-10-14
PRIOR APPLICATION NUMBER: 60/283,794
PRIOR FILING DATE: 2001-04-13
; PRIOR FILING DATE: 2001-04-13
; PRIOR APPLICATION NUMBER: 60/303,689
PRIOR FILING DATE: 2001-07-06
; NUMBER OF SEQ ID NOS: 70
; SOFTMARE: Patentin Ver: 2.1
; SEQ ID NO 69
LENGTH: 118
TYDE: DETERMATE
Search completed: December 17, 2004, 18:38:03 Job time: 230.978 secs
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 68
LENGTH: 118
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                                                                                                                                                                                                                                                                                Query Match 77.6%;
Best Local Similarity 81.5%;
Matches 97; Conservative 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: polypeptide
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ORGANISM: Artificial Sequence
FEATURE:
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                                                                                         60 LDSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCTRGFGDGGYFDVWGQGTLVTVSS 118
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                                                                                                                                                                                                                                                                                4;
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GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.

OM protein - protein search, using sw model

Run on: December 17, 2004, 18:11:22; Search time 46.7978 Seconds (without alignments) 244.665 Million cell updates/sec

Title: Perfect score: US-10-089-500-9
624
1 EVQLVESGGDFVQPGGSLRV.....KLGTYYFDSWGQGTLLTVSS 119

Sequence:

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched:

Total number of hits satisfying chosen parameters: 283416

283416 seqs, 96216763 residues

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR 79:\*
1: pir1:\*
2: pir2:\*
3: pir3:\*
4: pir4:\*

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

o . v 4	464.5		400	400	466	466	466	466.5	469	469.5	470	470.5	470.5	471	473.5	474	474	474	477.5	477.5	479	480	482.5	483	484	486	501	2	No. Score M	Result Qu
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Ig heavy chain V r	heavy chain -	heavy	neavy chain v	heavy chain v	heavy chain V	heavy chain V	Ig heavy chain V r	i-tetanus	heavy chair	heavy chain V-	heavy chain V	variak	heavy chain (	heavy chain pr	heavy chain V	heavy	heavy chain V	heavy chain	heavy chain pr	chain V	heavy	heavy chain V		heavy cha	Description					

45	44	43	42	41	40	39	38	37	36	35	34	<b>3</b> 3	32	<u>3</u> 1	30
453.5	453.5	453.5	454.5	455	457	457.5	458.5	459	459.5	461.5	461.5	462	462.5	463.5	463.5
72.7	72.7	72.7	72.8	72.9	73.2	73.3	73.5	73.6	73.6	74.0	74.0	74.0	74.1	74.3	74.3
133	122	118	120	119	123	122	141	140	114	137	136	139	128	147	114
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831510	S20772	S31116	S31112	B34353	S26794	E36005	S31669	A30532	S46390	S78054	S31587	137781	S26786	137780	S31120
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## ALIGNMENTS

RESULT 2 S31666 Ig heavy chain V region - human (fragment) C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Accession: S31666 R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C. submitted to the EMBL Data Library, June 1992 A;Description: Mechanisms that generate human immunoglobulin diversity operate from the A;Reference number: S31585 A;Accession: S31666 A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-138 <cui> A;Cross-references: EMBL:Z14202; NID:g30963; PIDN:CAA78571.1; PID:g30964 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin homology <imm></imm></cui>	Qy 61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS 119	OY 1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY 60	Query Match 80.3%; Score 501; DB 2; Length 119; Best Local Similarity 79.8%; Pred. No. 3.7e-38; Matches 95; Conservative 8; Mismatches 16; Indels 0; Gaps 0;	RESULT 1  Spacies: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999 C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999 C;Accession: S31107 R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, Eur. J. Immunol. 22, 247-251, 1992 A;Title: Restricted utilization of germ-line V(H)3 genes and short diverse third compleme A;Reference number: S31104; MUID:92111633; PMID:1730252 A;Accession: S31107 A;Accession: S31107 A;Accession: S31107 A;Accession: Fallor, mucleic acid sequence not shown; translation not shown A;Accession: S1107 A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Cross-references: EMBL:X62955 A;Cross-references: EMBL:X62955 A;Cross-references: EMBL:X62955 A;Rote: the nucleotide sequence was submitted to the EMBL Data Library, October 1991 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin homology <imm></imm>
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C.Species: Homo sapiens (man)
C.Dec: 02-Dec: 1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
C.Accession: S31108
R.Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman Bur. J. Immunol. 22, 247-251, 1992
A.Title: Restricted utilization of germ-line V(H)3 genes and short diverse third complem A.Reference number: S31104; MUID:9211633; PMID:1730252
A.Accession: S31108
A.Status: preliminary; nucleic acid sequence not shown; translation not shown
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A;Gene: GDB:IGH@; IGHDY1
A;Cross-references: GDB:118731; OMIM:146910
A;Map position: 14q32.33-14q32.33
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 16-Dec-1998
C;Accession: D36005
R;Schroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy chain variable A;Reference number: A36005; MUID:90349571; PMID:2117273
A;Accession: D36005
                                                                                                  A;Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1991 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <IMM>
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A; Residues: 1-119 < RAA>
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A; Residues: 1-119 < SCH>
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77.3%;
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                                                 77.68;
                            78.2%;
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Score 484; DB 2;
Pred. No. 1.2e-36;
9; Mismatches 17
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Pred. No. 8.2e-37;
9; Mismatches 18
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                                                 Length 119;
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Ig heavy chain precursor V region (mu) - human (fragment) C;Species: Homo sapiens (man) C;Jate: 24-Uul-1998 #sequence_revision 24-Uul-1998 #text_change C;Jate: 24-Uul-1998 #sequence_revision 24-Uul-1998 #text_change C;Accession: $70442 R;Cuisinier, A.M.; Fumoux, F.; Fougereau, M.; Tonnelle, C. Mol. Immunol. 29, 1363-1373, 1992 A;Title: IgM kappa/lambda EBV human B cell clone: an early step A;Reference number: $70442; MUID:93024508; PMID:1383695 A;Accession: $70442
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IS 1686
IS 1686
IS 1686
IS 1687
C; Species: Homo sapiens (man)
C; Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C; Accession: S31686
R; Couisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
submitted to the EMBL Data Library, June 1992
A; Description: Mechanisms that generate human immunoglobulin diversity operate from the
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A;Cross-references: EMBL:Z14205; NID:g30969; PIDN:CAA78574.1; PID:g30970 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;34-117/Domain: immunoglobulin homology <IMM>
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C;Superfamily: immunoglobulin V re
F;34-117/Domain: immunoglobulin ho
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Pred. No. 1.8e-36;
R; Mismatches 17;
                                                                                                                                               Score 482.5; DB
Pred. No. 2e-36;
9; Mismatches
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ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDHIVGATYFDYWGQGTLVTVSS 139

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A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-125 <MAR>
A;Cross-references: UNIPR
C;Superfamily: immunoglob
C;Keywords: heterotetrame
F;15-98/Domain: immunoglc
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A;Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1991
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995
C;Accession: S30531
R;Mariette, X.
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A;Title: Restricted utilization of germ-line V(H)3 genes and short diverse third complem A;Reference number: S31104; MUID:92111633; PMID:1730252
A;Accession: S31114
A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type; mRNA
A;Residues: 1-123 <RAAP
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C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
C;Accession: S31114
R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              submitted to the EMBL Data Library, October 1992 A; Reference number: $30520
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75.6%;
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Pred. No. 2.9e-36;
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Ig heavy chain V region - human (;Species: Homo sapiens (man) (;Species: Homo sapiens (man) ()Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 23-Mar-2001 (;Accession: S20782 R.Mortari, F.; Wang, J.; Schroeder, H.W. submitted to the EMBL Data Library, April 1992 submitted to the EMBL Data Library, April 1992 A,Description: Analysis of the IgA and IgG rearranged VH repertoire of huma A;Reference number: S20765
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A;Residues: 1-124 <MOR>
A;Residues: 1-124 <MOR>
A;Cross-references: EMBL:Z11946; NID:g33897; PIDN:CAA78003.1; PID:g33898
A;Cross-references: EMBL:Z11946; NID:g33897; PIDN:CAA78003.1; PID:g33898
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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Eur. J. Immunol. 22, 241-245, 1992
A;Title: The human cord blood antibody repertoire. Frequent usage of the V(H)7 gene famil
A;Reference number: S26786; MVID:92111632; PMID:1730251
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A; Residues: 1-128 < MOR>
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                                                                                                                                                    61 ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTÄVYYCAKERIAIFGVVIPHFDYWGQGTLV 120
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GTLVTVSS 128
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Ig heavy chain V region - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Date: 52-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31598
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
submitted to the EMBL Data Library, June 1992
A,Description: Mechanisms_that generate human immunoglobulin diversity oper.
  Ig heavy chain V region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31588
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C. submitted to the EMBL Data Library, June 1992
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S31588
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Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990

A;Title: Preferential utilization of conserved immunoglobulin heavy

A;Reference number: A36005; MUID:90349571; PMID:2117273
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C;Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 16-Aug-1996
C;Accession: C36005
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C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin C;Keywords: heterotetramer; immunoglobulin homology <IMM>
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A; Residues: 1-119 < SCH>
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ilarity 75.6%;
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Gauthier, L.; Boo
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77.3%;
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Pred. No. 1.1e-35;
9; Mismatches 20;
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Pred. No. 9.8e-36;
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A;Description: Mechanisme
A;Reference number: S3158
A;Accession: S31588
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-140 <CUI>
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A; Rolecule type: mRNA
A; Residues: 1-120 <MAH>
A; Residues: 1-120 <MAH>
A; Cross-references: EMBL: Z46382; NID: g562324; PIDN: CAA86521.1; PID: g1340167
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 15-98/Domain: immunoglobulin homology < IMM>
                                                                                                                     C;Species: Homo sapiens (man)
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change
C;Accession: S05271; S04602
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               A; Molecule type: mRNA
A; Residues: 1-160 <KIS1>
                                                                       submitted to the EMBL Data Library, A; Reference number: S05270
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                                                                                                                                                                                 Ig heavy chain precursor - human (fragment)
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Pred. No. 1.2e-35;
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R;Kishimoto, T.; Okajima, H.; Okumoto, T.; Taniguchi, M.
Nucleic Acids Res. 17, 4385, 1989
A;Title: Nucleotide sequences of the cDNAs encoding the V-regions of H-
A;Reference number: 804601; MUID:89296497; PMID:2500644
A;Accession: 804602
A;Molecule type: mRNA
A;Residues: 1-144 <KIS2>
A;Cross-references: EMBL:X14584
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;1-19/Domain: signal sequence #status predicted <SIG>
F;20-160/Product: Ig heavy chain (fragment) #status predicted <MAT>
F;34-117/Domain: immunoglobulin homology <IMM>
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A;Title: Restricted utilization of germ-line V(H)3 genes and short divers A;Reference number: $31104; MUID:92111633; PMID:1730252
A;Accession: $31105
A;Status: preliminary; nucleic acid sequence not shown; translation not say, Notecule type: mENA
A;Residues: 1-118 <a href="Residues">Residues</a>: 1-118 <a href="Residues">Residues</a
C;Species: Homo sapiens (man)
C;Date: 16-Feb-1996 #sequence_revision 16-Feb-1996 #text_change 23-Jul-1999
C;Accession: 13778; S25472
R;Demaison, C.; Chastagner, P.; Theze, J.; Zouali, M.
Proc. Natl. Acad. Sci. U.S.A. 91, 514-518, 1994
A;Title: Somatic diversification in the heavy chain variable region genes e
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C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 26-May-1995 #text_change 23-Jul-1999
C;Accession: S31105
R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurmar
Eur. J. Immunol. 22, 247-251, 1992
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Pred. No. 2.5e-35;
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Ig heavy chain V region (M49) - human (C.Species: Homo sapiens (man) C.Species: Homo sapiens (man) C.Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_chang C.Accession: F36005 R.Schroeder Jr., H.W.; Wang, J.Y. R.Schroeder Jr., H.W.; Wang, J.Y. Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990 A./Title: Preferential utilization of conserved immunoglobulin A./Reference number: A36005; MUID:90349571; PMID:2117273
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A;Accession: I37778
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-135 <RES>
A;Cross-references: EMBL:X67909; NID:g33574; PIDN:CAA48107.1; PID:g33575
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;28-111/Domain: immunoglobulin homology <IMM>
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C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin C;Keywords: heterotetramer; immunoglobulin homology <IMM>
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76.5%;
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Pred. No. 2.3e-35;
                                                                                                                                                                                                             Score 470; DB 2;
Pred. No. 2.2e-35;
                                                                                                                                                                                            Mismatches
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Ig heavy chain V-D-J region - human (Species: Homo sapiens (man) (C;Species: Homo sapiens (man) (C;Date: 13-Jan-1995 #sequence\_revision 13-Jan-1995 #text\_change 2 (C;Accession: S44111 R;Hawkins, R.E.; Zhu, D.; Ovecka, M.; Winter, G.; Hamblin, T.J.; submitted to the EMBL Data Library, March 1994 A;Description: Idiotypic vaccination against human B-cell lymphom A;Reference number: S44105

human B-cell lymphoma: rescue of variable

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24-May-2001

Stevenson,

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RESULT 21
PC2398
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A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-120 <HAW>
A;Residues: 1-120 <HAW>
A;Cross-references: EMBL:Z31387; NID:g472965; PIDN:CAA83262.1; PID:g940522
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
A;Cross-references: EMBL:X75388
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;15-98/Domain: immunoglobulin homology <IMM>
                                                                                                         Gene 148, 167-168, 1994
A;Title: Phage display of a human antibody against Clostridium tetani toxin.
A;Reference number: PC2397; MUID:95011651; PMID:7926831
A;Accession: PC2398
                                                                                                                                                                                                              anti-tetanus toxin 9F12 Fab heavy-chain - human (fragment) C;Species: Homo sapiens (man) C;Date: 13-Jun-1995 #sequence_revision 14-Jul-1995 #text_cC;Accession: PC2398
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S38489
                                                                  A; Molecule type: mRNA
A; Residues: 1-122 < ESP>
                                                                                                                                                                       R; Esposito, G.; Scarselli, E.; Traboni, Gene 148, 167-168, 1994
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A;Description: Human antibody fragments specific for human blood group antigens from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ig heavy chain - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
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A; Residues: 1-127 < MAR>
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A;Accession: S38489
A;Status: preliminary
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71.7%;
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Pred. No. 2.5e-35;
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Pred. No. 2.9e-35;
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Ig heavy chain V region - human C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 03-Mar-1994 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999 C;Accession: S31603 R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C. submitted to the EMBL Data Library, June 1992 A;Description: Mcchanisms that generate human immunoglobulin diversity oper
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S31603
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A;Map position: 14q32.33-14q32.33
C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <IMM>
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Proc. Natl. Acad. Sci. U.S.A. 37, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin
A;Reference number: A36005; MUID:90349571; PMID:2117273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ig heavy chain V region (M74) - human
C;Species: Homo sapiens (man)
C;Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 09-Jul-2004
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A; Residues: 1-121 < SCH>
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A; Status: preliminary
                   A; Accession: S31603
                                      A; Reference number: S31585
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73.8%;
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Pred. No. 4.7e-35;
2; Mismatches 17
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Pred. No. 5.2e-35;
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A; Molecule type: mRNA
A; Residues: 1-134 <CUI>A; Residues: 1-134 <CUI>CUIS: Seferences: EMBL: Z14203; NID: 930965; PIDN: CAA78572.1; PID: 930966
A; Cross-references: EMBL: Z14203; NID: 930965; PIDN: CAA78572.1; PID: 930966
C; Superfamily: immunoglobulin vegiobulin homology
C; Keywords: heterotetramer; immunoglobulin critical control of the control 
                                                                                                                                                                                                                                                                                                                                                                                                         C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999 (;Accession: S31679)
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C. submitted to the EMBL Data Library, June 1992
A;Description: Mechanisms that generate human immunoglobulin diversity oper. A;Reference number: S31585
A;Accession: S31679
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A;Title: Genetic analysis of self-associating immunoglobulin G rheumatoid factors A;Reference number: S23623; MUID:92156804; PMID:1740665
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A;Cross-references: EMBL:Z14168; NID:g30999; PIDN:CAA78537.1; PID:g31000 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
                                                                                                            C;Keywords: neceroceccame, F;34-117/Domain: immunoglobulin homology
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A; Residues: 1-143 <OLE>
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C;Species: Homo sapiens (man)
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C;Species: Homo sapiens (man)
C;Date: 13-Jan-1995 #sequence_revision 13-
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Best Local S
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76.5%;
                                          74.5%;
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; Pred. No. 6.2e-35;
11; Mismatches 15
       6,
Score 465; DB 2;
Pred. No. 7.1e-35;
6; Mismatches 17
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Pred. No. 5.7e-35;
7; Mismatches 19
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                                                                         Length 134;
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A;Residues: 1-139 <CUI>A;Residues: 1-139 <CUI>A;COSS-references: EMBL:Z14204; NID:g30967; PIDN:CAA78573.1; PID:g30968 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin C;Keywords: heterotetramer; immunoglobulin homology <IMM>
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C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31674
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
submitted to the EMBL Data Library, June 1992
A;Description: Mechanisms that generate human immunoglobulin diversity operate from
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C;Superfamily: immunoglobulin V region;
F;15-98/Domain: immunoglobulin homology
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A;Cross-references: GDB:120086; OMIM:147020
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C;Genetics:
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A; Residues: 1-121 < RES>
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A;Accession: 155673
A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Knight, G.B.; Agnello, V.; Bonagura, V.; Barnes, J.L.; Panka, J. Exp. Med. 178, 1903-1911, 1993
A;Title: Human rheumatoid factor cross-idiotypes. IV. Studies on tinct from the 17.109 and G6 XIds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ig heavy chain - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 21-Jan-2000
C;Accession: I55673
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A; Accession: S31674
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        EVQLLESGGGLVQPGGSLRLSCTASGFTFSTYGMSWVRQAPGKGLEWVSAISGSGGSTYY
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                                                                                                                                                  Score 464; DB 2; L
Pred. No. 7.9e-35;
Pred. No. 7.9e-35;
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Pred. No. 8.2e-35;
9; Mismatches 19
                                                                                                                                                                                                                                                                                                                    immunoglobulin homology
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Ig heavy chain V region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31699
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C. submitted to the EMBL Data Library, June 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT
S19666
                                                                                                                                                                       A; Molecule type: mRNA
A; Residues: 1-134 <CUI>
A; Cross-references: EMBL: Z14201; NID:g30961; PIDN:CAA78570.1; PID:g30962
A; Cross-references: EMBL: Z14201; NID:g30961; immunoglobulin homology
C; Superfamily: immunoglobulin v region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 34-117/Domain: immunoglobulin homology <IMM>
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A; Accession: S31699
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SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVSS
                                              EVQLLESGGGLVHPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYY
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                                                                                                                        74.4%;
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76.08;
                                                                                                                                                                                                                                                                                                               generate human immunoglobulin diversity operate from
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                                                                                                          Score 464; DB 2;
Pred. No. 8.8e-35;
7; Mismatches 17
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Pred. No. 7.9e-35;
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                                                                                                                                        Length 134;
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S26786
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RESULT 30

S31120
Ig heavy chain - human
Ig heavy chain - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999
C;Accession: S31120
C;Accession: S31120
R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, M.J.H.; Van Tol, M.J.D.; Vossen, J.M.; Schuurman, R;Raaphorst, M.J.H.; Van Tol, M.J.D.; Vossen, J.M.; Schuurman, M.J.H.; Van Tol, M.J.D.; Vossen, J.M.; Schuurman, M.J.H.; Van Tol, M.J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: EMBL:X67943; NID:g33578; PIDN:CAA48130.1; PID:g33579 C;Superfamily: immunoglobulin V region; immunoglobulin homology F;28-111/Domain: immunoglobulin homology <IMM>
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A;Title: Somatic diversification in the heavy chain variable A;Reference number: A36876; MUID:94119917; PMID:8290556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ig variable region (VDJ) (clone T20-11) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 16-Feb-1996 #sequence_revision 16-Feb-1996 #text_change
C;Accession: I37780; S25474
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A,Title: Restricted utilization of germ-line V(H)3 genes
A,Reference number: S31104; MUID:92111633; PMID:1730252
A,Accession: S31120
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A; Residues: 1-147 < RES>
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A; Residues: 1-114 < RAA>
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                                                                                                                                                                                                                                                                   61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTR--VKLGTYYF---DSWGQGTLL 115 : | | | | | | | | | | | | : | 115
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93;
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                                                                                                                                                                                                                                 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKDGEGWGLYYYYGMDVWGQGTTV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     llarity 75.0%;
Conservative
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74.8%; Pred. No. 8.2e-35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      74.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 463.5; DB 2
Pred. No. 1.1e-34;
7; Mismatches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
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Ig heavy chain V region - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change
C;Accession: S26786
R;Mortari, F; Newton, J.A.; Wang, J.Y.; Schroeder Jr., H.W.
Eur. J. Immunol. 22, 241-245, 1992
A;Title: The human cord blood antibody repertoire. Frequent usa;
A;Reference number: S26786; MUID:92111632; PMID:1730251
A;Accession: S26786
A;Status: preliminary
A;Molecule type: mENA
A;Residues: 1-128 <MOR>
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31587
C;Accession: A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
                                                                                                                                                     RESULT 34
S31587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: EMBL:X67908; NID:g33580; PIDN:CAA48106.1; PID:g33581
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;35-118/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Accession: I37781
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-139 <RES>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Species: Homo sapiens (man)
C;Date: 16-Feb-1996 #sequence_revision 16-Feb-1996 #text_chan
C;Accession: I37781; S25475
C;Accession: I37781; S25475
R;Demaison, C; Chastagner, P.; Theze, J.; Zouali, M.
Proc. Natl. Acad. Sci. U.S.A. 91, 514-518, 1994
A;Title: Somatic diversification in the heavy chain variable
A;Reference number: A36876; MUID:94119917; PMID:8290556
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                                                                                                       Ig heavy chain V region - human
C;Species: Homo sapiens (man)
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Best Local S
Matches 89
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Best Local &
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                                                                                                                                                                                                                                         VDSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCARKRAGKAAFNIWGQGTMVTVSS 139
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69.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                          9,
                                                                                                                                                                                                                                                                                                                                                                                                                 Score 462; DB 2;
Pred. No. 1.4e-34;
9; Mismatches 21
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Pred. No. 1.1e-34;
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A;Description: Mechanisms that generate human immunoglobulin diversity operate from the A;Reference number: S31585
A;Recession: S31587
A;Rotesul: preliminary
A;Molecule type: mRNA
A;Residues: 1-136 <CUI>A;Cross-references: EMBL:Z14189; NID:g31005; PIDN:CAA78558.1; PID:g31006
C;Keywords: heterotetramer; immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology <IMM>
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                                                                                                                                                                                                                                              F;1-i9/Domain: signal sequence #status predicted <SIG>F;20-137/Product: Ig heavy chain (fragment) #status pr F;34-116/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                          A;Molecule type: mRNA
A;Residues: 20-116 <HAW>
A;Cross-references: EMBL:X54435
C;Superfamily: immunoglobulin V region; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: EMBL:X54435; NID:g37812; PIDN:CAA38304.1; R;Harindranath, N.; Goldfarb, I.S.; Ikematsu, H.; Burastero, SInt. Immunol. 3, 865-875, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Species: Homo sapiens (man)
C;Date: 19-Nov-1997 #sequence revision 05-Dec-1997 #text_change 23-Jul-1999
C;Accession: S78054; S23719
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A; Residues: 1-137 < HAR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       submitted to the EMBL Data Library, August 1990 A;Reference number: S78051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R; Harindranath,
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                                                                                                                                                                   Matches
                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                             Keywords: immunoglobulin
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                                                                                                          1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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92; Conserv
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EVQLVESGGGLVQPGGSLRLSCVVSGFPFSTYWMHWVRQTPGKGLVWVAMINSGGTGT-Y
                                                                                                                                                                   Conservative
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76.7%;
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76.5%;
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                                                                                                                                                                   9; Mismatches
                                                                                                                                                                                  Score 461.5; DB 2;
Pred. No. 1.5e-34;
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Pred. No. 1.5e-34;
7; Mismatches 20;
                                                                                                                                                                                                                                                                     predicted <MAT>
                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
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                                                                                                                                                                                                       Length 137;
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Ig heavy chain V region - human C;Species: Homo sapiens (man) C;Date: 27-Jan-1995 #sequence\_revision 27-Jan-1995 #text\_change C;Accession: S46390 R;Figini, M.; Marks, J.D.; Winter, G.; Griffiths, A.D. J. Mol. Biol. 239, 68-78, 1994

20-Jun-2000

RESULT 36

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A; Introns:
C; Superfami
C; Keywords:
F; 20-140/Pr
F; 34-117/Do
                                                      Ig heavy chain V region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31669
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C. submitted to the EMBL Data Library, June 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          J. Immunol. 141, 3994-3998, 1988
A,Title: Lack of detectable somatic hypermutation in the V region A;Reference number: A30532; MUID:89035542; PMID:3141510
A;Accession: A30532
A;Accession: Lack of Accessions A;Residues: 1-140 <MEE>
A;Cross = references: GB:M23110
C;Genetics:
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C;Date: 31-Dec-1989 #sequence_revision 31-Dec-1989 #text_change 18-Oct-1996
C;Accession: A30532
R;Meeker, T.C.; Grimaldi, J.C.; O'Rourke, R.; Loeb, J.; Juliusson, G.; Einh J. Immunol: 141, 3994-3998, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Title: In vitro assembly of repertoires of antibody chains on the surface A;Reference number: S46390; MUID:94254092; PMID:8196048 A;Accession: S46390 A;Status: preliminary A;Molecule type: DNA A;Residues: 1-114 <FIG- A;Cross-references: EMBL:231686; NID:g509782; PIDN:CAA83491.1; PID:g1335143 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin C;Keywords: heterotetramer; immunoglobulin P;15-98/Domain: immunoglobulin homology <IMM>
A; Reference number: S31585
A; Accession: S31669
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                                        A; Description: Mechanisms that
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Best Local S
Matches 92
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Best Local
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20-140/Product: Ig heavy chain V-III region 38 #status predicted <MAT>
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                                        generate human immunoglobulin diversity operate from
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Pred. No. 2.6e-34;
2; Mismatches 15
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Pred. No. 1.9e-34;
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A;Status: prelimina
A;Molecule type: mRNA
A;Residues: 1-141 <CUI>
A;Cross-references: EMBL:Z14212; NID:g30959; PIDN:CAA78581.1; PID:g30960
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
F;34-117/Domain: immunoglobulin homology <IMM>
F;34-117/Domain: immunoglobulin homology <IMM>
Ig heavy chain V region - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 17-Mar-1999
C;Accession: S26794
R;Mortari, F:, Newton, J.A.; Wang, J.Y.; Schroeder Jr., H.W.
Eur. J. Immunol. 22, 241-245, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Gene: GDB:IGH@; IGHDY1
A;Gene: GDB:IGH@; IGHDY1
A;Cross-references: GDB:118731; OMIM:146910
A;Map position: 14q32.33-14q32.33
C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Schroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy chain variable gene :
A;Reference number: A36005, MUID:90349571; PMID:2117273
A;Accession: E36005
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A; Residues: 1-122 < SCH>
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Pred. No. 2.9e-34;
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A;Title: The human cord blood antibody repertoire. Frequent usage of the V(H)7 gene fami A;Reference number: $26786; MUID:92111632; PMID:1730251
A;Recession: $26794
A;Status: preliminary
A;Molecule type: mRNA
A;Residuse; l-123 <MOR>
A;Cross-references: EMBL:X61011
C;Superfamily: immunoglobulin v region; immunoglobulin homology
C;Reywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IPMN>
Query Match
Best Local Similarity 73.2%; Score 457; DB 2; Length 123;
Best Local Similarity 73.2%; Pred. No. 3.4e-34;
Matches 90; Conservative 11; Mismatches 18; Indels 4; Gaps 2;
Qy
1 EVQLIVESGGDIVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGSGTYY 60
Db 1 EVQLIVESGGGLVQPGGSLRVSCAASGFTFSSYSMYWVRQAPGKGLEWVSYISSSSTIYY 60
Qy
61 SDSVKGRFTISRDNAKNSLYLQMNSLRDEDTAVYYCARSIKYYDENYYGMDVWGQGTTVT 120
Qy
117 VSS 119
Db 121 VSS 119
Search completed: December 17, 2004, 18:30:18
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: uniprot_sprot:*
2: uniprot_trembl:*
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AAH18747
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AAL35865
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AAH24289
AAL35877
BAD00534
Q6MZU6
          Q9UL91
BAD00233
Q6GMY2
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Bad00459 camelis d

Q6mzq6 homo sapien

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Q6gmx2 homo sapien

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Adhla747 homo sapien
Bad00491 camelus d
O8wukl homo sapien
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O6pi81 homo sapien
Adh1376 homo sapien
Adh13965 lama glam
O9ul91 homo sapien
Bad00233 camelus d
O6gmy2 homo sapien
Bad00448 camelus d
Bad00510 camelus d
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Bad00444 camelus d
Bad00444 camelus d
Aah24289 homo sapi
Aal35877 lama glam
Bad00534 camelus d
Q6mzu6 homo sapien
Cae45931 homo sapien
                                                               Q9ul72 homo sapien
Bad00459 camelus d
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RESULT 2
BAD00406
ID BAD0
AC BAD0
DT 02-M

BAD00406 PRELIMINARY; PRT; BAD00406; 02-MAR-2004 (TrEMBLrel. 27, Created)

128 AA.

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			Query Best I Matche	PRC NON SEC	HSS Int Int	fet You	Wu DEC	Hom Euk Man NCB	MY 0 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	RESULT 1 Q9UL71			2 C C C C C C C C C C C C C C C C C C C	
119 S 119     121 S 121	61 SDSYKGRFTISRDNSKNTLYLØMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVS    -	1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYIS 	Query Match 75.2%; Score 469; DB 2; Length 1 Best Local Similarity 74.4%; Pred. No. 2.4e-41; Matches 90; Conservative 11; Mismatches 18; Indels	IGV; 1. 5; IG_LIKE; 1. 121 121 13, 13154 MW; 2F045CCFA5D50736 CRC64	AF035043, AAD56279.1; AF035043, AAD56279.1; P01852; 1NFD. P01852; 1NFD. P118007110; Ig-like. P118003596; Ig_v. P118003596; Ig_v. P118003596; Ig_v. P118003596; Ig_v.	D.C.; n-reactive autoantibodies n; n;	L1) SEQUENCE FROM N.A. SEQUENCE FROM N.A. MEDIJINE=98277139; PubMed=9614934; Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney	agment). o sapiens (Human). aryota; Metazoa; Chordata; Craniata; Vertebrata; malia; Eutheria; Primates; Catarrhini; Hominidae; I_TaxID=9606;	(TrembLrel. 13, Created) (TrembLrel. 13, Last sequence update) (TrembLrel. 25, Last annotation update) tive immunoglobulin heavy chain variable	ULT 1 L71 Q9UL71 PRELIMINARY; PRT; 121 AA.	ALIGNMENTS	442.5 70.9 125 2 BAD00494 442.5 70.9 131 2 BAD00494 442.5 70.9 465 2 Q6P6C4 442.5 70.9 465 2 AAH62335 441.5 70.8 127 2 BAD00445 441.7 70.7 124 2 BAD00513	444 71.2 122 2 BAD00446 444 71.2 122 2 BAD00549 443.5 71.1 118 2 AAL35882 443.5 71.1 129 2 BAD00424 443.5 71.1 473 2 Q6MZV7 443.5 71.1 473 2 CAE45950 443.5 71.1 473 2 CAE45950 443.5 71.1 473 2 CAE45950	
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Submitted (SEP-2002) to the EMBL/GenBank/DDBJ
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Mammalia; Eutheria;
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Submitted (SEP-2002) to the EMBL/GenBank/DDBJ
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Mammalia; Eutheria;
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02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
Immunoglobulin heavy chain VHDJ region (Fragment).
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"Libraries of heavy-chain antibodies reflecting camel gamma2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Spleen;
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                                        DSVKGRFTISRDNSKNTLYLOMRSLRAEDSAVYFCTRVK-----LGTYYFDSWGQGTLL
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                                                                                     VQLVESGGGLVQPGGSLRLSCAASGFTFSSYDMSWVRQAPGKGLEWVSAINSGGGSTYYA
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Lin heavy chain VHDJ region (Fragment).
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Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
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Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
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71.0%;
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Pred. No. 2.3e-40;
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Pred. No. 5.4e-41;
LOMNSLKPEDTAVYYCATREYGGSWLVGYYSMDYWGKGTLV
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WEDLINE-2238057; PubMed=12477932;

A Klausner R.D., Cellins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

Klausner R.D., Cellins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Lischul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Lischul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Wockernan K.J., Malek J.A., Gunaratne P.H.,

A Bosak S.A., McEwan P.J., McKernan R.J., Lu X., Gibbs R.A.,

A Hilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

Xones S. T. Marra M.A.
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Best Local :
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02-MAR-2004 (TrEMBLrel.
02-MAR-2004 (TrEMBLrel.
02-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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Mammalia; Eutheria;
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Homo sapiens (Human)
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Primates;
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73.8%;
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Pred. No. 1.:
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Catarrhini;
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AAL35877 AAL35877; 02-MAR-2004

PRELIMINARY;

117

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RESULT 6
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Best Local Similarity
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MEDLINE=22007448; PubMed=12009207;

MEDLINE=22007448; PubMed=12009207;

Tanha J., Dubuc G., Hirama T., Narang S.A., MacKenzie C
"Selection by phage display of llama conventional V(H)
heavy chain antibody V(H)H properties.";

J. Immunol. Methods 263:97-109(2002).

EMBL; AF442946; AAL35877.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Libraries of heavy-Chain antibodies reflecting gamma3 in vivo repertoires.";
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NON_TER
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Cetartiodactyla; Tylopoda; Camelidae; Lama.
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Mammalia; Eutheria;
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02-MAR-2004
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121
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117 AA;
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(TrEMBLrel. 27, Last sequence update)
(TrEMBLrel. 27, Last annotation update)
lin heavy chain VHDJ region (Fragment).
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(TrEMBLrel. 27, Last annotation update)
in heavy chain variable domain (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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12459 MW;
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Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
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73.1%;
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Pfam; PF00047; 19; 4.

SMART; SM00409; IG; 2.

SMART; SM00409; IGc1; 3.

SMART; SM00406; IGv; 1.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG LIKE; 4.

PROSITE; PS00290; IG_MHC; UNKNOWN_2.

Hypothetical protein.

SEQUENCE 464 AA; 51099 MW; 2FCA72C
                  TISSUE-Human rectum tumor;
Bloecker H., Boecher M., Mewe
Fobo G., Han M., Wiemann S.;
Submitted (AUG-2003) to the E
EMBL; BX640874; CAB45931.1; -
Hypothetical protein.
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02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation updat
Hypothetical protein DKFZp686C15213.
DKFZP686C15213.
                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primata; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                        CAE45931;
02-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                     CAE45931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bloecker H., Boecher M., Mewes H.W., Weil B., Amid C., (Fobo G., Han M., Wiemann S.; Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases EMBL; BX640874; CAE45931.1; -.
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hypothetical protein DKFZp686C15213.
Name=DKFZp686C15213;
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    SEQUENCE
                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                               Homo sapiens (Human)
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                                                                                                                                                                                             NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                     PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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Primates;
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    51099
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75.0%;
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27, Last sequence up
27, Last annotation
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Pred. No. 2.7e-39;
3; Mismatches 15
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Catarrhini;
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i; Hominidae; Homo.
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                                                                                                                                                            A Strauberg R.;

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Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases.

R EMBL; BC018747; AAH18747.1; -.

R InterPro; IPR003599; Ig.

R InterPro; IPR003597; Ig-q1.

R InterPro; IPR003597; Ig-q1.

R InterPro; IPR003596; Ig-wc.

R Ffam; PF07654; C1-Bet; 3.

R Pfam; PF07654; C1-Bet; 3.

R SMART; SM00409; IG; 2.

R SMART; SM00409; IG; 2.

R PROSITE; PS00290; IG LIKE; 4.

PROSITE; PS00290; IG LIKE; 4.

PROSITE; PS00290; IG-MHC; UNKNOWN_2.
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A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
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A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,
A Scapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
A Scapleton M., Gares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
A Scapleton M., Gares M.B., Bonaldo M.F., Mullahy S.J.,
A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
A Robak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
A Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gubbs R.A.,
A Hithard S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
A Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
A Wilting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
A Whiting M., Touchman J.W., Green B.D., Dickson M.C.,
A Hones S. T. May Touchman J.W., Green B.D., Dickson M.C.,
A Kozywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
A Jones S. T. Marra M. A.
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Matches 90; Conserv
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Mammalia; Eutheria;
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05-JUL-2004
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05-JUL-2004
                                                                                                                            Hypothetical protein seQUENCE 470 AA;
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MEDLINE=22388257; PubMed=12477932;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Generation and initial analysis of more than 15,000 full-length humar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mouse cDNA sequences.";
c. Natl. Acad. Sci. U.S.A.
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Last sequence up
Score 456; DB 2;
Pred. No. 2.7e-39;
7; Mismatches 22;
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Pred. No. 2.7e-39;
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                                                            Length 470;
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REDLINE-2238857; PubMed-12477932;

REDLINE-2238857; PubMed-12477932;

REDLINE-2238857; PubMed-12477932;

RIAGRAM ALBERT R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RIAGRAM ALBERT R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.,

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RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Wadin T.B., Foarcia A.M., Gau M., Gunaratne P.H.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Pahey J., Helton E., Ketteman M., Madan A., Gay L.J., Hulyk S.W.,

RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Ra Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Ra Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

RA Kones S. T. Marra M.A.
                                                                                                                                                                                                                                                                                                                                         Query Match
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Matches 90
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Homo sapiens (Human).
Eukaryota; Metazoa; Ch
Mammalia; Eutheria; Pı
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02-MAR-2004 (TrEMBLrel.
02-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Strausberg R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Jones S.J., Marra M.A.; "Generation and initial analysis of more than 15,000 full-length and mouse cDNA sequences.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hypothetical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Primary B-Cells;
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Last annotation update)
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7; Mismatches 22;
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Best Local
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            TISSUE=Primary B-Cell8;

MEDLINE=2238857; PubMed=12477932;

Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunrattne P.H.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren E.J., LX., Glibbs R.A.,

Villalon D.K., Muzny D.M., Sodergren E.J., LX., Glibbs R.A.,
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Q8WUK1;
01-MAR-2002
01-MAR-2002
01-MAR-2004
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BAD00491;
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Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases
EMBL; AB092129; BAD00491.1; -.
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Mammalia; Eutheria;
NCBI_TaxID=9606;
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Homo sapiens (Human)
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Mammalia; Eutheria;
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les of heavy-chain antibodies reflecting
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  Helton E.,
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(TrEMBLrel. 27, Last sequence update)
(TrEMBLrel. 27, Last annotation update)
in heavy chain VHDJ region (Fragment).
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13593 MW;
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Primates;
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Chordata, Craniata, Vertebrata, Euteleostomi,
Cetartiodactyla, Tylopoda, Camelidae, Camelus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      73.0%;
  Ketteman M.,
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); Mismatches
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  Rodrigues S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     update)
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Sanchez A.,
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Q9UL90
ID Q9UL9
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AC Q1LMA
DT 01-MA
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DT 01-MA
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CFrag
COS MAMMA
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Best Local
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InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003597; Ig_MHC.
InterPro; IPR003596; Ig_V.
Pfam; PF007654; Cl-set; 4.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGV; 1.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS50835; IG_MHC; UNKNOWSEQUENCE 613 AA; 67295 MM; 6
                    fetus.";
Clin. Im
                                                                                                                                                                                                                                                                                                                                                                                                    Q9UL90;
                                                                                                                                                                                                                                                                                                          01-WAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
Wu X., Liu B., Van der Merwe P.L.,
                                                                                                                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jones S.J., Marra M.A.;
"Generation and initial analysis
and mouse cDNA sequences.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffan
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Bu
Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A.,
                                                                                                                                                                                                                                                                     Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (DEC-2001) to the EMBL; BC020240; AAH20240.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A. TISSUE-Primary B-C
                                                                  "Myosin-reactive autoantibodies in rheumatic
                                                                                          Young
                                                                                                                                                                                                                                                                                         (Fragment)
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PH1642

PH1642

PH1643

PH1645

PH1645

PH1645

PH1645

PH1646

PH1098

PL0120

PL0120

PL0120

PL0120

S11590

S11590

S11590

S31116

S31116

S31119

S31119

S31119
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                                                                                          D.C.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTR-VKLGTYYFDSWGQGTLLTVSS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYDGSNKYY
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                                                                                                                                                                                                                         Chordata;
Primates;
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75.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 455.5; DB 2
Pred. No. 4.2e-39;
                                                                                                                                                                                                                      Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99:16899-16903 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               60C7F5950671E315 CRC64;
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                                                                                                              Kalis N.N.,
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                                                                     carditis
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                                                                                                              Berney
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Schein J.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     613;
                                                                       normal
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Immunol. Immunopathol. AF035024; AAD56260.1;

87:184-192 (1998)

S78486; S78486.

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Q 968BB9
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                                                                                                                                                                                                                                                                                                                                                                                                                        XX MEDILIE=2388257; Pubmed=12477932;

XX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

XX Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

XX Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

XX Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

XX Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

XX Altschenko L., Marusinia K., Farmer A.A., Rubin G.M., Hong L.,

XX Diatchenko L., Marusinia K., Farmer A.A., Rubin G.M., Hong L.,

XX Altschenko L., Marusinia K., Bonaldo M.F., Casavant T.L., Schetz T.E.,

XX Altschenko L., Marusinia K.B., Bonaldo M.F., Casavant T.L., Schetz T.E.,

XX Altschenko L., Marusinia K.B., Bonaldo M.F., Casavant T.L., Schetz T.E.,

XX Altschenko L., Wagner M.B., Bonaldo M.F., Casavant T.L., Schetz T.E.,

XX Altschenko L., Wagner M.B., Bonaldo M.F., Casavant T.L., Schenez T.F.,

XX Altschenko L., Marusinia K.J., Malek J.A., Gunaratte P.H.,

XX Altschenko L., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

XX Altschenko L., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

XX Altschenko L., Skalska U., Shevchenko Y., Bouffard G.G.,

XX Altschenko L., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

XX Altschenko L., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

XX Altschenko L., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

XX Altschenko L., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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Best Local S
Matches 89
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InterPro; IPR007110; IG-
InterPro; IPR003596; Ig-
InterPro; IPR003596; Ig-
Pfam; PF00047; 19; 1.
SMART; SM00406; IGV; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q96BB9;
Q96BB9;
01-DEC-2001
01-DEC-2001
01-MAR-2004
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       InterPro;
InterPro;
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                                                                                                                                                                              Submitted (OCT-2001) to 1
EMBL; BC015760; AAH15760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IGHM protein.
Homo sapiens
                                                                              InterPro;
                                                                                                                                                                                                                                  Strausberg R.
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                                                         InterPro;
                                                                                                          HSSP;
                                                                                                                                                                                                                                                         FISSUE=Primary
                                                                                                                                                                                                                                                                                                                                                                                      "Generation
                                                                                                                                                                                                                                                                                                                                                                                                           Jones S.J., Marra M.A.;
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                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14
                                                    ; S05271; S05271.
; S24260; S24260.
P; P01861; 1ADQ.
erPro; IPR007110;
erPro; IPR003597;
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                               IPR003006;
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                                                                                                                                                                                                                                                                                                                               and initial analysis DNA sequences."; Acad. Sci. U.S.A. 99:
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                                                                                                                                                                                                                                                            B-Cells;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B-Cells;
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12437 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chordata;
Ig-like.
Ig_cl.
Ig_MHC.
Ig_v.
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                                                                                                                                                                                                                                                                                                                                  U.S.A. 99:16899-16903(2002)
                                                                                                                                                                                the 0.1;
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Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Created)
                                                                                                                                                                                                     EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 455; DB 2;
Pred. No. 6.7e-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., Ugdin T.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Nillalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Hopky J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Holton S., Woung A.C., Shevchenko Y., Bouffard G.G.,
RA Holton S., Carimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Holton S. M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Holton S. M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Holton S. M. J., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Holton S. M. J., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Holton S. M. J., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Holton S. M. J., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Holton S. M. J., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Holton S. M. J., Skalska U., Smailus D.E., Schein J.E.,
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Best Local S
Matches 90
                         Submitted (DEC-2002) to the EMB EMBL; BC041037; AAH41037.1; -. InterPro; IPR003599; Ig. like. InterPro; IPR0037110; Ig-like. InterPro; IPR003597; Ig.cl. InterPro; IPR003596; Ig_MHC. InterPro; IPR003596; Ig_WHC. InterPro; IPR003596; Ig_WHC. InterPro; IPR003596; Ig_WHC.
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05-JUL-2004
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Pfam; PF00047; 19; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS00290; IG_MHC; UNKNOWN
SEQUENCE 597 AA; 65039 MW; 4FC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hypothetical protein. Homo sapiens (Human).
                                                                                                                                                                                                                                                                       Strausberg R
                                                                                                                                                                                                                                                                                                                                                                                                                           "Generation and initial analysis of more and mouse cDNA sequences.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=22388257; PubMed=12477932;
Strausberg R.L., Feingold E.A., Grouse L.H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q6PI81;
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                                                                                                                                                                                                                                                                                                          TISSUE=Primary
                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jones S.J., Marra M.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15
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                                                                                                                                                                                                                                                                                                          B-Cells;
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Primates;
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Last sequence update)
Last annotation updat
                                                                                                                                                                                                                                                 EMBL/GenBank/DDBJ databases
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Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                     99:16899-16903 (2002)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
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4FCA3AD8ECE263D9 CRC64;
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les 17;
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AC AAH411
AC AAH411
AC AAH411
AC AAH41
DT 02-MA
DT 102-MA
DT 102-
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RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

RA Hopkins R.F., Jordan H., Moore T., Roandan T.L., Scheetz T.E.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Raha S.S., MocEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Hichards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Hopkinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Rodriguez R.C., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

RA Rodriguez R.G., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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SMART; SM00407; IGG1; 3.
SMART; SM00406; IGG; 1.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS00290; IG_MHC; UNKNOWN_2.
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AAH41037;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Primary B-Cells;
MEDLINE=22388257; PubMed=12477932;
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hypothetical protein.
Homo sapiens (Human).
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                                                                                                                   Hypothetical protein. SEQUENCE 478 AA; 5
                                                                                                                                                                         Submitted (DEC-2002) to the EMBL; BC041037; AAH41037.1;
                                                                                                                                                                                                                                                                                                                                               and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.
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                                                                                                                                                                                                                            Strausberg R.;
                                                                                                                                                                                                                                                            FISSUE=Primary
                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Jones S.J., Marra M.A.;
                                                                                                                                                                                                                                                                                                                                                                                                        'Generation and initial analysis of more than 15,000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          QGTLLTVSS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VDSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCAREFESTMTTVNADYYYFYMDVWG
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     Conservative
                                                                                                                                                                                                                                                            B-Cells;
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Primates;
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71.3%;
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71.3%;
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Last annotation update)
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                                                                                                                                                                                                    EMBL/GenBank/DDBJ databases
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     7;
                                                           Score 454;
Score 454; DB 2; I
Pred. No. 4.5e-39;
7; Mismatches 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                                                   17BED38D917970D6 CRC64;
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4.5e-39;
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                                                        Length 478;
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Best Local S
Matches 87
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Q9UL91;
01-MAY-2000
                        fetus.";
Clin. In
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SEQUENCE FROM N.A.
MEDLINE-98277139; PubMed=9614934;
MEDLINE-98277139; Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MBL; AF NON_TER
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=22007448; PubMed=12009207;
Tanha J., Dubuc G., Hirama T., Narang S.A.,
"Selection by phage display of llama convenu
heavy chain antibody V(H)H properties.";
J. Immunol, Methods 263:97-109(2002).
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Tylopoda; Camelidae; Lama.
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                                                                                                                                                                                                        NCBI_TaxID=9606;
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Mammalia; Eutheria;
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02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
Immunoglobulin heavy chain variable domain (Fragment).
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                                                                         "Myosin-reactive autoantibodies in rheumatic
                                                                                                                                                                                                                                                                                   Homo sapiens (Human).
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                                                                                                                                                                                                                                                                                                         (Fragment).
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Immunol. Immunopathol. AF035023; AAD56259.1;
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73.1%;
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Primates;
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Pred. No. 1.2e-39;
                             87:184-192(1998)
                                                                                                                                                                                                                                       Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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                                                                                                                     Kalis N.N.,
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02-MAR-2004
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 Q6GMY2;
05-JUL-2004
                                                                                                                                                                                                                                                                                                                               TISSUE-Peripheral blood;
Honda T., Akahori Y., Kurosawa Y.;
Honda T., Akahori Y., Kurosawa Y.;
"Libraries of heavy-chain antibodies reflecting camel gamma2
gamma3 in vivo repertoires.";
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig-v.
Pfam; PP00047; Ig; 1.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PIR;
PIR;
PIR;
                          Q6GMY2
                                                                                                                                                                                                                                                                                                 NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9838;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Camelus dromedarius
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                        EMBL; AB091871; BAD00233.1; -.
                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunoglobulin heavy
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S21205; S21205.
S30531; S30531.
                                                                                      121
                                                                                                            116 TVSS 119
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                                                                                                                                      61
                                                                                                                                                   88;
                                                                                                                                                                                                  1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                     TISS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SDSVKGRFTISRDNSKNTLYLOMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTLLTVS 118
                                                                                                                                      ADSVKGRFTISRDNAKNTLYLQLNSLKTEDTAMYYCAKDTVVAAPKWDYSMDYWGKGTLV
                                                                                                                                                                                    QVQLQESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAINSGGGSTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYSMNWVRQAPGKGLEWVSYISSTIITIYY
                                                                                                                                                                                                                                                                                    124
124 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          118 AA; 12843 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (TrEMBLrel. 27, Created)
(TrEMBLrel. 27, Last sequence update)
(TrEMBLrel. 27, Last annotation update)
in heavy chain VHDJ region (Fragment).
                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRELIMINARY;
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(TrEMBLrel. 27,
                         PRELIMINARY;
                                                                                      124
                                                                                                                                                                                                                                                                                                                                                                                                                                  (Dromedary) (Arabian camel).
Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
                                                                                                                                                                                                                                               72.4%;
68.5%;
                                                                                                                                                                                                                                                                                     13374 MW;
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                                                                                                                                                                                                                                    15;
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 Created)
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5; Mismatches
                                                                                                                                                                                                                                            Score 451.5; DB 2
Pred. No. 1.7e-39;
                         PRT;
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                                                                                                                                                                                                                                                                                     FE75B41506CA8D25 CRC64;
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BAD00448 ID BAD0 AC BAD0

BAD00448; BAD00448;

PRELIMINARY;

PRT;

125 AA

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RX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RX Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Brownstein M.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Brownstein M.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA RA S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA RICHARDS S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Rabtey J., Helton E., Ketteman M., Wadan A., Rodrigues S., Sanchez A.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

Jones S.J., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human

"Generation and initial analysis of more than 15,000 full-length human

"Generation and initial analysis of more than 15,000 full-length human

"Generation and Sci. U.S.A. 99:16899-16903 (2002).
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Best Local S
Matches 90
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InterPro; IPR003599; IG-like.

InterPro; IPR007110; Ig-like.

InterPro; IPR0031597; Ig_cl.

InterPro; IPR003006; Ig_MHC.

InterPro; IPR003006; Ig_MHC.

InterPro; IPR003906; Ig_V.

Pfam; PF007654; Cl-set; 4.

Pfam; PF00047; ig; 4.

SMART; SM00400; IG; 2.

SMART; SM00400; IGc1; 4.

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05-JUL-2004 (TrEMBLrel. 2
Hypothetical protein.
Homo sapiens (Human).
Bukaryota; Metazoa; Chord
Mammalia; Eutheria; Prima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Strausberg R.;
Submitted (JUN-2004) to the
EMBL; BC073758; AAH73758.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical protein. SEQUENCE 606 AA; 6
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                                                                                                                                                                             107
                                                                                                                                                                                                                                                                                                      61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTR------
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                                                                                                                                                                                                                                                                                                                                                                                       90;
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                                                                                                                                                                                                                                                   ADSVKGRFTISRDNAKNSLYLOMNSLRAEDTAVYYCARGGNGIAAAGRVVYAEDYYYYYG
                                                                                                                                                                         -DSWGQGTLLTVSS 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   72.4%; Score 451.5;
67.2%; Pred. No. 1.1e
tive 12; Mismatches
                                                                                                          153
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27,
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Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .1e-38;
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RESULT 22
BAD00510
ID BAD000
AC BAD00
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DT 02-MA
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Best Local S
Matches 85
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Best Local S
Matches 87
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02-MAR-2004
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Mammalia; Eutheria;
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02-MAR-2004
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Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases
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"Libraries of heavy-chain antibodies reflecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted
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02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
10-minoglobulin heavy chain VHDJ region (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Libraries of heavy-chain antibodies reflecting camel gamma2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Honda T., Akahori Y.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SSUE=Spleen;
                                                                                                                                                                                                                                                                                                                                                                                                                                    3 in vivo repertoires.";
tted (SEP-2002) to the EMBL/GenBank/DDBJ
AB092148; BAD00510.1; -.
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                                                                                             EVQLVESGGGLVQPGGSLRLSCAASAITFSSYSMTWVRQAPGKGLEWVSVISSSGDSTYY
                                                                                                                                                                                                                                                                                                                                              126
126 AA;
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(TrEMBLrel. 27, Last sequence update)
(TrEMBLrel. 27, Last annotation update)
                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cetartiodactyla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (Dromedary) (Arabian camel).
Chordata; Craniata; Vertebrata; Eutel.
Cetartiodactyla; Tylopoda; Camelidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13671 MW;
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Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
                                                                                                                                                                                                                                                                                                                                              13665 MW;
                                                                                                                                                                                                                                                    72.0%;
69.1%;
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71.3%;
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                                                                                                                                                                                                                                                    Score 449; DB 2;
Pred. No. 3.2e-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 449.5;
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RESULT 24
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Best Local
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NON_TER
SEQUENCE
Honda T., Akahori Y., Kurosawa Y.; "Libraries of heavy-chain antibodies gamma3 in vivo repertoires."; Submitted (SEP-2002) to the EMBL/GenE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Young D.
                                                                                                                                                                                                                                                             IGVH.
                                                                                                                                                                                                                                                                                02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
102-MAR-2004 (TrEMBLrel. 27, Last annotation update)
1mmunoglobulin heavy chain VHDJ region (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-CCT-2003 (TrEMBLrel. 25, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
                                                                                     TISSUE=Spleen;
                                                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                         Camelus dromedarius
                                                                                                                                                                                                                                                                                                                                                                               BAD00459
                                                                                                                                                                                                                                                                                                                                                                                                        BAD00459
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SMART; SM00406; IGv; 1.
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PIR; S21205; S21205.
HSSP; P01783; 11GC.
InterPro; IPR007110; Ig-like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
Wu X., Liu B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens (Human)
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01-MAY-2000
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                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           fetus."
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                                                                                                                                                                   NCBI_TaxID=9838;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROSITE; PS50835; IG_LIKE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EVQLVESGGGLVQPGGSLRLSCAASGFTVSSNYMNWVRQAPGKGLESVSVTYSGGS-SYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                        PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       118
12872 MW; B4D1A5944B2D5CCA CRC64;
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Primates;
                                                                                                                                                                                         (Dromedary) (Arabian camel).
Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    71.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10;
    EMBL/GenBank/DDBJ databases
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Pred. No. 3.4e-39;
0; Mismatches 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Matches 85
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InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig-dl.
InterPro; IPR003597; Ig-dl.
InterPro; IPR003596; Ig-MC.
InterPro; IPR003596; Ig-v.
Pfam; PP007654; C1-set; 3.
Pfam; PP00767; ig; 4.
SMART; SM00409; IG; 2.
SMART; SM00407; IGc1; 3.
SMART; SM00407; IGC1; 3.
SMART; SM00406; IGv; 1.
PR0SITE; PS00290; IG_MHC; UNKNOWN_2.
Hypothetical protein.
SEQUENCE 475 AA; 52043 MW; B7EAE25
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SEQUENCE
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Lauber J., Bahr A., Mewes H.W., Weil B., Amid
Han M., Wiemann S.;
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ
EMBL; BX640947; CAE45972.1; -.
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein DKFZp686G11190.
Name=DKFZp686G11190;
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                              EVQLLESGGGLVQPGGSLRLSCAASGFTFRNYAMNWVRQAPGKGLEWVSGISSSGVNTYY
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12957 MW; A55F627A3F977A6D CRC64;
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Pred. No. 1.7e-38;
3; Mismatches 19
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Pred. No. 3.5e-39;
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3; Mismatches
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                                                                                                 AMEDINE-22388557; PubMed=12477932;

Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Histchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brah S.S., Loquellano N.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Wagner R.J., Malek J.A., Gunaratne P.H.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Raha S.S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hullyk S.W.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hullyk S.W.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Schein J., Kanghan J.W., Green E.J., Lu X., Gibbs R.A.,

A Willialon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Pahey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

A Whiting M., Madan A., Young A.C., Shevchenko Y., Boutfard G.G.,

A Holder M. J., Skalska U., Smailus D.E., Schnerch A., Schein J.S.,

A Torog G. T. M. Salska U., Smailus D.E., Schnerch A., Schein J.S.,
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05-JUL-2004 (TrEMBLrel. 2
05-JUL-2004 (TrEMBLrel. 2
Hypothetical protein.
Homo sapiens (Human).
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Hypothetical protein.
Hypothetical protein.
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02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein DKFZp686G11190.
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                                    Jones S.J., Marra M.A.; "Generation and initial analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE=Spleen,
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Lauber J., Bahr A., Mewes H.W., Weil
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Mammalia; Eutheria;
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87; Conserv
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Primata; (
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Primates;
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. 27, Last sequ
. 27, Last anno
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Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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; Hominidae;
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Matches 89; Conserv
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Interpro; IPR003597; Ig.cl.
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Interpro; IPR003596; Ig.MHC.
Interpro; IPR003596; Ig.v.
Pfam; PF00647; Ag.; 3.
Pfam; PF00047; Ag.; 3.
SMART; SM00409; IG.; 4.
SMART; SM00407; IGC1; 2.
SMART; SM00408; IGV; IG.
PR0SITE; PS50835; IG.LIKE; 4.
PR0SITE; PS50835; IG.MHC; UNKNOWN 1.
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02-MAR-2004
02-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
TISSUE=Peripheral blood;
Honda T., Akahori Y., Kurosawa Y.;
"Libraries of heavy-chain_" reflecting camel gamma2
                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                        NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gamma3 in vivo repertoires.";
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ
EMBL; AB091863; BAD00225.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BAD00225;
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SEQUENCE 49
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   SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKL--GTYY----FDSWGQGT
                                                                            QVQLVQSGGGLVQPWGSLRLSCAASGFTFSGYAMSWVRQAPGKGLEWVSAINSGGGSTYY
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126 AA;
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(TREMBLrel. 27, Last sequence update)
(TREMBLel. 27, Last annotation update)
Lin heavy chain VHDJ region (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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13717 MW; 486E1D741474EF26 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (Dromedary) (Arabian camel).
Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                52865 MW; 55B999305B286203 CRC64;
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73.6%;
                                                                                                                                                                                                                 71.6%; Score 446.5; DB 2
68.3%; Pred. No. 5.9e-39;
tive 17; Mismatches 16
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Pred. No. 2.6e-38;
9; Mismatches 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99:16899-16903 (2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 126
                                                                                                                                                                                                                                                                                          DB 2; Length 126;
                                                                                                                                                                                                                         16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   databases
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                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Camelus
                                                                                                                                                                                                                     7;
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                                                                                                                                                                                                                         Gaps
                                                                                                                                                    6
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                                                                            60
                                                                                                                                                                                                                         <u>د</u>
RESULT 30
BAD00420
ID BAD00
AC                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT
Q920ET
ID Q920ET
AC Q920ET
DT 001
DT 
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Query Match
Best Local S
Matches 89
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Q1-DEC-2001
Q1-DEC-2001
Q1-OCT-2003
                                                                                                                                                                                                  02-MAR-2004
02-MAR-2004
02-MAR-2004
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(Fragment).
(Mouse).
Mus musculus (Mouse).
Metazoa; Chordata; (
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Rodentia;
                                                                                                                                                                                                                                                   BAD00420
BAD00420;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00047; ig; 1.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Atkin J.D., Iape A., Jennings
Submitted (SEP-2000) to the EN
EMBL, AF307937; AAL09421.1; -.
            TISSUE-Spleen;
Honda T., Akahori Y., Kurosawa Y.;
"Libraries of heavy-chain antibodies reflecting
                                                                                            NCBI_TaxID=9838;
                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                Camelus dromedarius
                                                                                                                                                                              02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation updat
Immunoglobulin heavy chain VHDJ region (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PIR; C25913; C25913.
HSSP; P01783; 1IGC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=10090;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q920E7
                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   121
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                                                                                                                                                                                                                                                                                                                                                                                    119 S 119
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                                                                                                                                                                                                                                                                                                                                                                                                                                            61 SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTY--YFDSWGQGTLLTVS
                                                                                                                                                                                                                                                                                                                                                                                                                       61 PDSVKGRFTISRDNAKNTLYLQMSSLKSEDTAMYYCAR--HGDYDVGFAYWGQGTLVTVS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EVOLVESGGDLVKPGGSLKLSCAASGFTFSSYGMSWVRQTPDKRLEWVATISSGGSYTYY
VIVO
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119 AA;
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(TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      71.5%; Score 446; DB 2; L
73.6%; Pred. No. 6.3e-39;
73.6%; Original Press. No. 6.3e-39;
                                                                                                             (Dromedary) (Arabian camel).
Chordata; Craniata; Vertebrata;
Cetartiodactyla; Tylopoda; Camel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13025 MW; F6E904044381CA7C CRC64;

    Created)
    Last sequence update)
    Last annotation update)

                                                                                                                                                                                 VHDJ region (Fragment).
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EMBL/GenBank/DDBJ da
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                     PRT;
                                                                                                                                                                                                                                                                     126
                                                                                                                                                                                                    ion update)
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                                                                                                                  Camelidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 119;
                 camel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                  Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               region
                 gamma2
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               and
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RESULT 32
BAD00446
ID BAD00
AC BAD00
DT 02-MAD0T 02-MADDT 02-MADDT 1GVH.
CS Camel
OC Mamma
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BAD00525
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Best Local S
Matches 85
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Best Local S
Matches 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (SEP-2
EMBL; AB092058; I
NON_TER 1
NON_TER 126
                                                     BAD00446;
BAD00446;
02-MAR-2004
02-MAR-2004
02-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BAD00525;
02-MAR-2004
02-MAR-2004
02-MAR-2004
                                                                                                                                                                                                                                                                                              NON_TER
  Eukaryota;
Mammalia;
                                                                                                                                                                                                                                                                                                                   gamma3 in vivo repertoires.";
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ
EMBL; AB092163; BAD00525.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                 Camelus dromedarius
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                       Camelus dromedarius
                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Spleen;
                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BAD00525
                                          02-MAR-2004 (TrEMBLrel. 27,
02-MAR-2004 (TrEMBLrel. 27,
02-MAR-2004 (TrEMBLrel. 27,
Immunoglobulin heavy chain V
                                                                                                                                                                                                                                                                                                                                                                                                                                                    IGVH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          [mmunoglobulin
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                                                                                                                                                                                                                                          85;
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                                                                                                                                                                                                                                                                                                                                                  ., Akahori Y., Kurosawa Y.;
ies of heavy-chain antibodies reflecting camel
                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                              VTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SDSVKGRFTISRDNSKNTLYLQMRSLRAEDSAVYFCTRVKLGTYYFDS------WGQGTL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QVQLVESGGGTVQPGGSLRLSCAASGFTFSTYAMSWVRQAPGKGLEWVSAINSGGGSTYY
                                                                                                                                                     SDSVKGRFTISRDNSKNTLYLOMRSLRAEDSAVYFCTRVKLGTYYFDSWGQGTILLTVSS 119
   Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SEP-2002) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              126 AA;
                                                                                                                                                                                                                                                                                     121 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          (TIEMBLrel. 27, Created)
(TIEMBLrel. 27, Last sequence update)
(TIEMBLEL. 27, Last annotation update)
lin heavy chain VHDJ region (Fragment).
                                                                                                  PRELIMINARY;
                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                 121
12951 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              126
13569 MW;
(Dromedary) (Arabian camel).
Chordata; Craniata; Vertebrata; Eutel
Cetartiodactyla; Tylopoda; Camelidae;
                                                                                                                                                                                                                                                                                                                                                                                                                 (Dromedary) (Arabian camel).
Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
                                                                                                                                                                                                                                                    71.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              71.5%;
                                            , Last
, Last
VHDJ r
                                                                                                                                                                                                                                          16;
                                                                            Created)
                                                                                                                                                                                                                                         Score 445.5; 1
Pred. No. 7.2e.
16; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL/GenBank/DDBJ
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Pred. No. 6.7e-39;
5; Mismatches 16
                                                                                                                                                                                                                                                                                     A6D50E79D505E5B4 CRC64;
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                                            region
                                          sequence update)
annotation update)
region (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 121
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                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                              Length
 Euteleostomi;
lidae; Camelus
                                                                                                                                                                                                                                                                                                                                                    gamma2
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                                                                                                                                                      118
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RASULT 33

BADOCS 10

AC BADOC

AC BADOC

AC BADOC

AC BADOC

DT 02-MA

DT 02-MA

DT 02-MA

DT 02-MA

CAMBLI

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RN 1
RESULT 34
AAL35882
ID AAL35
AC A-MA
DT 02-MA
DT 02-MA
DT 02-MA
DT 02-MA
DT 1mmun
OS Lama
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Best Local S
Matches 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 84
                                      AAL35882;
02-MAR-2004
02-MAR-2004
02-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; AF NON_TER NON_TER
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BAD00549;
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"Libraries of heavy-chain antibodies reflectir
gamma3 in vivo repertoires.";
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ
EMBL; AB092084; BAD00446.1; -.
NON TER 1 1
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                                                                                                                             AAL35882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gamma3 in vivo repertoires.";
Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Honda T., Akahori Y., Kurosawa Y.;
"Libraries of heavy-chain antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
TISSUE=Spleen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=9838;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Camelus dromedarius
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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TISSUE=Spleen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9838;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AB092187; BAD00549.1; -.
ER 1 1
ER 122 122
NCE 122 AA; 13354 NW;
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                                                                                                                                                                                                                                                                                                                                                                                                 83;
                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                               PDSVKGRFTISRDNAKNILYLQLNSLKIDDTATYYCAKEIGEDYCFDYWGQGTQVTVSS
                                                                                                                                                                                                                                                                                                                 QVQLVESGGGLVQPGGSI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Akahori Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    122
122 AA;
                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                    (TrEMBLrel.
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13395 MW;
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Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
                                                                                                                                                                                                                                                                                                                                                                                                                       71.2%;
69.7%;
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VHDJ region (Fragment).
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Pred. No. 1e-3
L4; Mismatches
                                                                                    Created)
                                                                                                                                                                                                                                                                                                                                                                                                                    Score 444; DB 2
Pred. No. 1e-38;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
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Immunoglobu

lin hea (Llama) (TrEMBLrel. (TrEMBLrel.

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Last sequence update)
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domain (Fragment).

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Best Local Similarity
Matches 87; Conserv
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Mammalia; Eutheri
NCBI_TaxID=9844;
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Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases.
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Honda T., Akahori Y., Kurosawa Y., "Libraries of heavy-chain antibodies
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129 AA;
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(TrENBLrel. 27, Last sequence update)
(TrENBLrel. 27, Last annotation update)
Lin heavy chain VHDJ region (Fragment).
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       PRELIMINARY;
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Chordata; Craniata; Vertebrata; Euteleostomi;
Cetartiodactyla; Tylopoda; Camelidae; Camelus
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73.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 443.5; DB 2;
Pred. No. 1.3e-38;
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Pred. No. 1.
       PRT;
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05-JUL-2004
05-JUL-2004
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02-MAR-2004 (TrEMBLrel.
02-MAR-2004 (TrEMBLrel.
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05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hypothetical pupier DKFZP686C11235
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02-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CAE45920
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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473 AA; 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (Human).
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Primates;
                              52121 MW;
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Pfam; PF07654; C1-set; 3.

Pfam; PF00047; 19; 4.

SMART; SM00409; IG; 2.

SMART; SM00400; IGC1; 3.

SMART; SM00406; IGV; 1.

PROSITE; PS50835; IG LIKE; 4.

PROSITE; PS508290; IG MHC; UNKNOWN 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE-Human small intestine;
Bloecker H., Boecher M., Mewes H.W., Weil B.,
Fobo G., Han M., Wiemann S.;
Submitted (AUG-2003) to the EMBL/GenBank/DDBJ
EMBL; Bx640853; CAE45920.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bloecker H., Boecher M., Mewes H.W., Weil B., Fobo G., Han M., Wiemann S.; Submitted (AUG-2003) to the EMBL/GenBank/DDBJ EMBL; BX640853; CAE45920.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=Human small intestine;
THE GERMAN HUMAN CDNA CONSORTIUM;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADSLOGRFTISRDNARNSLYLOMNSLRAEDTAVYYCARQNEHTSPWYPSFFDYWGQGILV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EIQLVESGGLVQPGGSLRLSCAASGFTFSSFEMNWVRQAPGKGLEWLSYITRSGNTVYY
                                                                                                                                                                                                                                                       (TrEMBLrel. 27, Last sequence up (TrEMBLrel. 27, Last annotation protein DKFZp686C11235.
                                                                                                                                                                                 Chordata; Craniata; Primata; Catarrhini;
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27, Last sequence update)
27, Last annotation update)
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                                                                                                                                                                                                                                                                                                                 Created)
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Pred. No. 5.7e-38;
8; Mismatches 18
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9476EAE4C0BFC447 CRC64;
                                                                                                                                                                                                                                                                                                                                                       473
                                                                                                                                                                                                   Vertebrata;
                                                                                                                                                                                     Hominidae;
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RESULT 39
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ID AAL35875;
AC AAL35875;
DT 02-MAR-2004 (
DE Immunoglobuli
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Q9UL93;
01-MAY-2000
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SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
MEDLINE=98277139; Van der Merwe P.L.
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PIR; PL0120; PL0120.
HSSP; P01772; 2FB4.
InterPro; IPR007110; I
02-MAR-2004 (TrEMBLrel. 27, Created)
02-MAR-2004 (TrEMBLrel. 27, Last sequence update)
02-MAR-2004 (TrEMBLrel. 27, Last annotation update)
Immunoglobulin heavy chain variable domain (Fragment).
Lama glama (Liama)
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Mammalia; Eutheria;
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01-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                  VQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVAYISSGGSGTYYS
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IPR003596; Ig_v.
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(TrEMBLrel. 13, Last sequence update)
(TrEMBLrel. 26, Last sequence update)
rive immunoglobulin heavy chain variable
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12434 MW;
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76.5%;
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Pred. No. 5.7e-38;
.8; Mismatches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 443; DB 2;
Pred. No. 1.3e-38;
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Catarrhini; Hominidae;
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02-MAR-2004 (TrEMBLrel. 27,
02-MAR-2004 (TrEMBLrel. 27,
01-mmunoglobulin heavy chain V
                                                                                                                                                                                                                                                                                                                                                                                                                                               gamma3 in vivo repertoires."; Submitted (SEP-2002) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TISSUE=Spleen;
Honda T., Akahori Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bukaryota; Metazoa;
Mammalia; Eutheria;
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MEDLINE=22007448; PubMed=12009207;
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Mammalia; Eutheria;
NCBI_TaxID=9844;
[1]
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                                                                                                                                                                                                      EVQLVESGGDFVQPGGSLRVSCAASGFAFSHYAMSWVRQAPGKGLEWVA---YISSGGSG
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122
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                                                                                                                                                                                                                                                                     12;
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, Last annotation update)
VHDJ region (Fragment)
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Pred. No. 1.5e-38;
2; Mismatches 21
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Pred. No. 1
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1.3e-38;
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